New River Basin

Cause Group Code: N01R-02-BAC Little Helton Creek

Cause Location: A tributary to Helton Creek. The segment extends from the Virginia state line upstream.

City / County: Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A Fecal Coliform / 5A

The ambient water quality monitoring station 9-LHC001.92 had a 75% exceedance of the E. coli water quality standard.

Recreation Greek Fecal Coliform - Total	lmn oire e	l Siza by Water Type	Estuary (Sq. Miles)		servoir .cres)	River (Miles) 6.30
VAS-N01R_LHC01A02 / Little Helton Creek & tributaries / From Virginia state line upstream to Haw Orchard in Grayson Highlands State Park, a tributary to Helton Creek, WQS Section 2, ii (NE02). Little Helton Creek	5A	Fecal Coliform		2004	M	6.30
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
Escherichia coli (E. coli) - Total	Impaired	Size by Water Type:				6.30
Little Helton Creek Recreation			Estuary (Sq. Miles)	Reservoir (Acres)		River (Miles
VAS-N01R_LHC01A02 / Little Helton Creek & tributaries / From Virginia state line upstream to Haw Orchard in Grayson Highlands State Park, a tributary to Helton Creek, WQS Section 2, ii (NE02).	5A	Escherichia coli (E. col)	2010	M	6.30
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size

Sources:

Grazing in Riparian or Shoreline Zones

Unrestricted Cattle Access

New River Basin

Cause Group Code: N02R-01-TEMP Wilson Creek

Cause Location: Wilson Creek mainstem from the New River confluence at Mouth of Wilson upstream 8.8 miles.

City / County: Grayson Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5A

Station 9-WLS001.78 had 2 of 11 temperature measurements exceed the water quality standard for Class VI waters.

Assessment Unit / Water Name / Location Desc.	Cause	e rry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N02R_WLS01A04 / Wilson Creek / Middle segment of Wilson Creek from mile 8.8 near Rugby, upstream to Quebec Branconfluence, WQS Section 2. Most is in National Forest (NE03).	5A ch	Temperature		2020	L	4.62
VAS-N02R_WLS01A98 / Wilson Creek / Wilson Creek mainster from New River confluence at Mouth of Wilson upstream 8.8 miles WQS Section 2. Parallel to Rt. 58, includes Volney (NE03).		Temperature		2020	L	8.90
Wilson Creek			Estuary	Res	servoir	River
Aquatic Life			(Sq. Miles)	(A	cres)	(Miles)
Temperature - Total	Impaire	d Size by Water Type:				13.52

Sources:

Source Unknown

New River Basin

Cause Group Code: N02R-02-BAC New River and Grassy Creek

Cause Location: This segment begins at the North Carolina state line, includes Fields Dam, and extends downstream to the New River confluence with Saddle Creek at the Route 601 bridge and from where the New River reenters VA from NC to the confluence with Peach Bottom Creek. Grassy Creek from the headwaters downstream to the North Carolina

state line and Bridle Creek, a tributary of the New River west of Rt. 601.

City / County: Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

Fecal Coliform / 5A

Priority

Size

0.73

Listed

2004

The AWQM station located at 9-NEW187.46 had a 42% exceedance of the E. coli water quality standard and station 9-NEW181.66 had a 18% exceedance. 9-NEW172.45 had 16% that exceeded WQS. Station 9-GRA003.36 had a 92% exceedance. Level III citizen data at station 9-BRL1-NCNR indicate a 46% exceedance rate and station 9-BRL000.04 had a 58% exceedance rate. Station 9-NEW171.94 had a 13% exceedance of the e.coli water quality standard.

Assessment Unit / Water Name / Location Desc.	Cause Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N02R_BRL01A10 / Bridle Creek / Tributary of New River, west of Rt. 601, south of Rt. 58 (NE07).	5A	Escherichia coli (E. coli)	2014	М	1.13
VAS-N02R_GRA01A10 / Grassy Creek / Headwaters to NC state line east of Quillen Ridge and parallel SR 725, WQS Section 2 (NE04).	5A	Escherichia coli (E. coli)	2010	M	3.64
VAS-N02R_NEW01A98 / New River / Upper mainstem begins at the North Carolina state line at river mile 189.06, and extends downstream to the Wilson Creek confluence at Mouth of Wilson at river mile 188.46. Headwaters are in North Carolina, WQS Section 2 (NE04)	5A	Escherichia coli (E. coli)	2010	M	0.73
VAS-N02R_NEW02A98 / New River / Mainstem from the Wilson Creek confluence downstream to the Fox Creek confluence near Fox, WQS Section 2k (NE04).	5A	Escherichia coli (E. coli)	2010	М	2.50
VAS-N02R_NEW03C04 / New River / Mainstem from Fox Creek confluence downstream to the Bridle Creek confluence at SR 601 bridge north of Big Ridge, WQS Section 2k (NE07).	5A	Escherichia coli (E. coli)	2010	M	4.22
VAS-N04R_NEW01A98 / New River / Mainstem from Brush Creek confluence downstream to Peach Bottom Creek confluence, parallel to North Carolina state line, WQS Section 2 (NE10)	(5A	Escherichia coli (E. coli)	2012	M	5.98
VAS-N04R_NEW01B02 / New River / New River mainstem north of Privett Knob, from Bridle Creek confluence downstream to Saddle Creek confluence, WQS Section 2 (NE08).	5A	Escherichia coli (E. coli)	2010	M	1.47
VAS-N04R_NEW02B06 / New River / From NC state line downstream to Brush Creek confluence at Rt. 21/221 bridge, WQS Section 2 (NE10).	5A	Escherichia coli (E. coli)	2020	L	0.42
New River and Grassy Creek			Estuary		servoir	River
Recreation Escherichia coli (E. coli). Total In	nnaira	d Sizo by Water Type:	(Sq. Miles) (A	cres)	(Miles)
Escherichia coli (E. coli) - Total In	ipaire	u Size by water Type:				20.09
	Caus	е		Cycle First	TMDL Dev.	Water

Category Cause Name

5A Fecal Coliform

Final 2020 Appendix 5 - 3384

Assessment Unit / Water Name / Location Desc.

the North Carolina state line at river mile 189.06, and extends

VAS-N02R_NEW01A98 / New River / Upper mainstem begins at

New River Basin							
	confluence at Mouth of Wilson at e in North Carolina, WQS Section						
VAS-N02R_NEW02A98 / New Creek confluence downstream to Fox, WQS Section 2k (NE04).	River / Mainstem from the Wilson of the Fox Creek confluence near	n 5A	Fecal Coliform		2004	M	2.50
VAS-N02R_NEW03C04 / New River / Mainstem from Fox Creek 5A Fecal Coliform confluence downstream to the Bridle Creek confluence at SR 601 bridge north of Big Ridge, WQS Section 2k (NE07).					2004	M	4.22
_	River / New River mainstem nort k confluence downstream to Sado 2 (NE08).		Fecal Coliform		2004	M	1.47
New River and Grassy Creek				Estuary	Res	ervoir	River
Recreation				(Sq. Miles)	(Ac	cres)	(Miles)
	Fecal Coliform - Total	Impaire	d Size by Water Type	э:			8.92
Sources:			·				
Grazing in Riparian or Shoreline Zones	Rural (Residential Areas)	Sourc	e Unknown	Unrestri	cted Ca	attle Acc	ess

New River Basin

Cause Group Code: N02R-02-HG New River

Cause Location: This segment begins at the upper mainstem at the North Carolina state line at river mile 189.06, and extends downstream to the Saddle Creek confluence, it includes the mainstem from the North Carolina line in N04

downstream to the confluence with Rock Creek and the mainstern from Buddle Branch downstream to the

confluence with Reed Creek.

City / County: Grayson Co. Wythe Co.

Use(s): Fish Consumption

Cause(s) / VA Category: Mercury in Fish Tissue / 5A

Station 9-NEW171.94 showed smallmouth bass, rock bass and carp exceeded the level of concern for Mercury; a second flathead catfish exceeded the Virginia Department of Health's level of concern. Station 9-NEW158.40 was monitored for sediment and fish tissue. Mercury exceeded the level of concern in several species. 9-NEW117.47 was monitored for sediment and fish tissue in 2004. Mercury was found in the fish tissue.

	Cause atego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N02R_NEW01A98 / New River / Upper mainstem begins at the North Carolina state line at river mile 189.06, and extends downstream to the Wilson Creek confluence at Mouth of Wilson at river mile 188.46. Headwaters are in North Carolina, WQS Section 2 (NE04)	5A	Mercury in Fish Tissue		2006	L	0.73
VAS-N02R_NEW02A98 / New River / Mainstem from the Wilson Creek confluence downstream to the Fox Creek confluence near Fox, WQS Section 2k (NE04).	5A	Mercury in Fish Tissue		2006	L	2.50
VAS-N02R_NEW03C04 / New River / Mainstem from Fox Creek confluence downstream to the Bridle Creek confluence at SR 601 bridge north of Big Ridge, WQS Section 2k (NE07).	5A	Mercury in Fish Tissue		2006	L	4.22
VAS-N04R_NEW01A98 / New River / Mainstem from Brush Creek confluence downstream to Peach Bottom Creek confluence, parallel to North Carolina state line, WQS Section 2 (NE10)	5A	Mercury in Fish Tissue		2006	L	5.98
VAS-N04R_NEW01B02 / New River / New River mainstem north of Privett Knob, from Bridle Creek confluence downstream to Saddle Creek confluence, WQS Section 2 (NE08).	5A	Mercury in Fish Tissue		2008	L	1.47
VAS-N04R_NEW01C02 / New River / Mainstem west of Baywood, from Little River confluence downstream to Rock Creek confluence, WQS Section 2 (NE12).	5A	Mercury in Fish Tissue		2006	L	4.68
VAS-N04R_NEW02A06 / New River / From Peach Bottom Creek confluence downstream to Little River confluence, WQS Section 2 (NE10).	5A	Mercury in Fish Tissue		2010	L	3.61
VAS-N04R_NEW02B06 / New River / From NC state line downstream to Brush Creek confluence at Rt. 21/221 bridge, WQS Section 2 (NE10).	5A	Mercury in Fish Tissue		2006	L	0.42
VAS-N08R_NEW03A06 / New River / Mainstem from I-77 bridge downstream to Reed Creek confluence near Lone Ash, WQS Section 2 (NE25).	5A 1	Mercury in Fish Tissue		2006	L	6.51
New River			Estuary (Sq. Miles)		servoir	River
Fish Consumption	naire	d Ciza by Water Trees	(3q. ivilles)	(A	cres)	(Miles)
Mercury in Fish Tissue - Total Im	paire	a size by water Type:				30.12

New River Basin

Sources:

Source Unknown

New River Basin

Cause Group Code: N02R-03-BAC Wilson Creek

Cause Location: This segment includes the Wilson Creek mainstem from the New River confluence upstream to the Quebec Branch

confluence.

City / County: Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A Fecal Coliform / 5A

The AWQM stations 9-WLS001.78 had a 30% and 9-WLS002.57 had a 25% exceedance of the E. coli water quality standard.

Assessment Unit / Water Name / Location Desc.	Cause Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N02R_LWL01A20 / Little Wilson Creek / Tributary to Wilson Creek, north of Mink Ridge, Section 2,ii (NE03).	5A	Escherichia coli (E. coli)		2020	L	7.57
VAS-N02R_WLS01A04 / Wilson Creek / Middle segment of Wilson Creek from mile 8.8 near Rugby, upstream to Quebec Branc confluence, WQS Section 2. Most is in National Forest (NE03).	5A ch	Escherichia coli (E. coli)		2018	M	4.62
VAS-N02R_WLS01A98 / Wilson Creek / Wilson Creek mainstem from New River confluence at Mouth of Wilson upstream 8.8 miles, WQS Section 2. Parallel to Rt. 58, includes Volney (NE03).	5A	Escherichia coli (E. coli)		2010	M	8.90
Wilson Creek			Estuary (Sq. Miles		servoir cres)	River (Miles)
Recreation Escherichia coli (E. coli) - Total II	mpaire	d Size by Water Type:	(Sq. Miles) (A	cies)	21.09
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N02R_WLS01A98 / Wilson Creek / Wilson Creek mainstem from New River confluence at Mouth of Wilson upstream 8.8 miles, WQS Section 2. Parallel to Rt. 58, includes Volney (NE03).	5A	Fecal Coliform		2004	М	8.90
Wilson Creek			Estuary		servoir	River
Recreation Fecal Coliform - Total II	mpaired	d Size by Water Type:	(Sq. Miles) (A	cres)	(Miles) 8.90

Sources:

Grazing in Riparian or Livestock (Grazing or Shoreline Zones Feeding Operations)

Source Unknown

New River Basin

Cause Group Code: N03R-01-BAC Fox Creek

Cause Location: This segment includes the mainstem of Fox Creek from the Mill Creek confluence to the New River confluence,
Middle Fox Creek from the Fox Creek confluence upstream 4.1 miles and Mill Creek from the confluence with Fox
Creek upstream to the headwaters. Little Fox Creek is included in this segment and it extends from the Fox Creek

confluence upstream 2.2 miles.

City / County: Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

Fecal Coliform / 5A

Designated a natural trout stream. The AWQM station, 9-FXC000.84 had a 33% exceedance of the e.coli water quality standard, 9-FXC003.35 had a 25%, station 9-LFX000.06 had a 45% exceedance of the E.coli standards, stations 9-MFX000.13 had a 50% exceedance and 9-MIR000.28 had a 33% exceedance of the E. coli water quality standard.

Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size	
VAS-N03R_FXC01A98 / Fox Creek / Mainstem of Fox Creek from Mill Creek confluence north of Grant to the New River confluence near Fox, WQS Section 2 (NE05)	5A	Escherichia coli (E. coli)		2010	M	7.65	
VAS-N03R_LFX01A10 / Little Fox Creek / A Fox Creek tributary downstream to confluence with Fox Creek, WQS Section 2, South of Grubbs Chapel, parallels Rt. 680 (NE06)	5A	Escherichia coli (E. coli)		2010	M	2.28	
VAS-N03R_MFX02A02 / Middle Fox Creek / From Fox Creek confluence upstream 4.4 miles, west of Buck Mountain, WQS Section 2, vi (NE06)	5A	Escherichia coli (E. coli)		2010	M	4.61	
VAS-N03R_MIR01A02 / Mill Creek / From Fox Creek confluence north of Grant, upstream to origin on Pine Mountain, WQS Section 2 ii, parallels Rt. 739 (NE05)	5A	Escherichia coli (E. coli)		2010	M	4.57	
Fox Creek			Estuary (Sq. Miles)		servoir .cres)	River (Miles)	
Recreation Escherichia coli (E. coli) - Total Im	npaired	d Size by Water Type:	(Sq. IVIIIes)	(,,	cres)	19.11	
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size	
VAS-N03R_FXC01A98 / Fox Creek / Mainstem of Fox Creek from Mill Creek confluence north of Grant to the New River confluence near Fox, WQS Section 2 (NE05)	5A	Fecal Coliform		2004	М	7.65	
Fox Creek			Estuary (Sa Miles)		servoir	River (Miles)	
Recreation (Sq. Miles) (Acres) Fecal Coliform - Total Impaired Size by Water Type:							

Sources:

Grazing in Riparian or Livestock (Grazing or Shoreline Zones Feeding Operations)

Source Unknown

New River Basin

Cause Group Code: N04R-02-BAC Little River

Cause Location: This segment includes the Little River mainstem from NC state line, river mile 5.20, to the confluence at New River.

City / County: Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The AWQM station, 9-LVR001.34, had a 25% exceedance of the fecal coliform water quality standard in the 2004 WQA. The station was moved to 9-NEW002.65 in 2003 and had a 16% exceedance of the E. coli water quality standard. Stations 9-LVR002.65 had a 25% exceedance and 9-LVR007.16 had a 28% exceedance of the E.coli water quality standard.

Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	Cyc Fir List	st Dev.	Water Size
VAS-N04R_LVR01A98 / Little River / Little River mainstem ear Peach Bottom, from NC state line, river mile 5.20, to the conflue of New River, WQS Section 2 (NE11)	•	. coli) 20°	12 M	6.55
Little River Recreation		Estuary (Sg. Miles)	Reservoir (Acres)	River (Miles)

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

6.55

Sources:

Rural (Residential Areas)

New River Basin

Cause Group Code: N04R-03-BAC Peach Bottom Creek and Rock Creek

Cause Location: This segment includes the mainstem from the headwaters downstream to the confluence with the New River. This also includes Rock Creek from the U.S. 21 crossing to the confluence with the New River.

City / County: Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

Fecal Coliform / 5A

The AWQM station, 9-PBC001.12, had a 83% exceedance of the E.coli water quality standard, at 9-PBC008.61, 58% exceed and at 9-RCK000.50, 100% exceed.

	Cause Catego	e rv Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N04R_PBC01A98 / Peach Bottom Creek / Mainstem from Beaverdam Creek confluence downstream to New River confluence parallel to SR 697, WQS Section 2 (NE09).	5A	Escherichia coli (E. coli)	1	2006	М	2.81
VAS-N04R_PBC01B02 / Peach Bottom Creek / Peach Bottom Creek headwaters north of Buck Mountain, downstream to confluence of Little Peach Bottom Creek north of Independence, WQS Section 2, vi (NE09).	5A	Escherichia coli (E. coli)	1	2012	М	8.86
VAS-N04R_PBC01C04 / Peach Bottom Creek / East of Independence from Beaverdam Creek confluence, upstream to Little Peach Bottom Creek confluence, WQS Section 2 (NE09).	5A	Escherichia coli (E. coli)	1	2016	М	5.34
VAS-N04R_RCK01A12 / Rock Creek / New River tributary from SR 654 near Chestnut Hill School downstream, northeast of Independence (NE12).	5A	Escherichia coli (E. coli)	1	2012	M	5.00
Peach Bottom Creek and Rock Creek Recreation			Estuary (Sq. Miles)		servoir cres)	River (Miles)
Escherichia coli (E. coli) - Total Im	npaired	d Size by Water Type:				22.01
	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N04R_PBC01A98 / Peach Bottom Creek / Mainstem from Beaverdam Creek confluence downstream to New River confluence parallel to SR 697, WQS Section 2 (NE09).	5A	Fecal Coliform		2004	M	2.81
Peach Bottom Creek and Rock Creek			Estuary (Sq. Miles)		servoir cres)	River (Miles)
Fecal Coliform - Total Im	npaired	d Size by Water Type:	(54. 14.1103)	(/-		2.81

Sources:

Source Unknown

New River Basin

Cause Group Code: N04R-07-BAC Saddle Creek

Cause Location: This segment includes the mainstem from the New River confluence upstream 3.09 miles, west of Independence.

City / County: Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The AWQM station, 9-SDL000.05, had a 42% exceedance of the E.coli water quality standard.

Cycle **TMDL** First Cause Dev. Water Priority Listed Size Assessment Unit / Water Name / Location Desc. Category Cause Name 2006 3.17 VAS-N04R_SDL01A06 / Saddle Creek / A New River tributary Escherichia coli (E. coli) west of Independence, WQS Section 2 (NE08)

Saddle Creek Estuary Reservoir River (Sq. Miles) (Acres) (Miles) Recreation Escherichia coli (E. coli) - Total Impaired Size by Water Type: 3.17

Sources:

Animal Feeding Operations Livestock (Grazing or (NPS)

Feeding Operations)

New River Basin

Cause Group Code: N04R-08-PH **New River**

Cause Location: The New River from the NC state line downstream to the Brush Creek confluence at the Rt. 21/221 bridge.

City / County: Grayson Co.

Use(s): Aquatic Life

Cause(s) / VA Category: pH / 5A

AWQM station 9-NEW171.94 had 2 of 18 pH measurements exceed the water quality standard.

Cycle **TMDL** First Cause Dev. Water **Priority** Category Cause Name Listed Size Assessment Unit / Water Name / Location Desc. 5A pH 2020 0.42 VAS-N04R_NEW02B06 / New River / From NC state line

downstream to Brush Creek confluence at Rt. 21/221 bridge, WQS

Section 2 (NE10).

New River Estuary Reservoir River (Sq. Miles) (Acres) (Miles) **Aquatic Life**

pH - Total Impaired Size by Water Type: 0.42

Sources:

Source Unknown

New River Basin

Cause Group Code: N05R-01-BAC Elk Creek & Tributaries

Cause Location: This segment includes Elk Creek from the Comers Rock Branch confluence downstream to the New River confluence, including 4.31 miles of Knob Fork and Middle Branch Elk Creek, west of Bennington Mill. It also

includes the headwaters of Turkey Fork near Dry Run Gap on Iron Mountain.

City / County: Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

Fecal Coliform / 4A

The AWQM stations, 9-EKC000.11, 9-EKC003.78, 9-EKC010.47, 9-EKC012.13, 9-EKC017.51 and 9-KNB000.03 had exceedances of the E. coli water quality standard that ranged from 26-66%. Stations 9-ECM001.01 had a 75% exceedance and 9-TKY001.55 had a 100% exceedance of the E. coli water quality standard.

·	Cause	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N05R_ECM01A14 / Middle Branch Elk Creek / Elk Creek tributary west of Bennington Mill, Section 2 (NE13).	4A	Escherichia coli (E. coli)		2016	L	3.06
VAS-N05R_EKC01A00 / Elk Creek / Lower Elk Creek from the Knob Fork confluence, north of Lundy Knob, downstream to the New River confluence, WQS Section 2,*** (NE14).	4A	Escherichia coli (E. coli)		2006	L	3.32
VAS-N05R_EKC02A00 / Elk Creek / Upper Elk Creek from the Turkey Fork confluence, north of Poor Knob, downstream to Knob Fork confluence, WQS Section 2, *** (NE14).	4A	Escherichia coli (E. coli)		2006	L	7.59
VAS-N05R_EKC03A02 / Elk Creek / Mainstem from confluence of Comers Rock Branch near Bennington Mill, downstream to Turkey Fork confluence, WQS Section 2 (NE13).	4A	Escherichia coli (E. coli)		2006	L	9.38
VAS-N05R_KNB01A06 / Knob Fork / Elk Creek tributary upstream to Farmers Branch, at The Pilot, NE of Brierpatch Mountain, WQS Section 2 (NE14).	4A	Escherichia coli (E. coli)		2006	L	4.60
VAS-N05R_TKY01A02 / Turkey Fork / Headwaters near Dry Run Gap on Iron Mountain in Jefferson National Forest, WQS Section 2 iii (NE14).	4A	Escherichia coli (E. coli)		2018	L	6.00
Elk Creek & Tributaries			Estuary		servoir	River
Recreation Escherichia coli (E. coli) - Total Imp	airec	d Size by Water Type:	(Sq. Miles)) (A	cres)	(Miles) 33.95
-	Cause atego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N05R_EKC01A00 / Elk Creek / Lower Elk Creek from the Knob Fork confluence, north of Lundy Knob, downstream to the New River confluence, WQS Section 2,*** (NE14).	4A	Fecal Coliform		2002	L	3.32
VAS-N05R_EKC02A00 / Elk Creek / Upper Elk Creek from the Turkey Fork confluence, north of Poor Knob, downstream to Knob Fork confluence, WQS Section 2, **** (NE14).	4A	Fecal Coliform		2002	L	7.59
Elk Creek & Tributaries			Estuary (Sq. Miles)		servoir cres)	River (Miles)
Recreation Fecal Coliform - Total Imp	aired	d Size by Water Type:	(=400)			10.91

New River Basin

Sources:

Animal Feeding Operations

(NPS)

Septage Disposal

Grazing in Riparian or Shoreline Zones

Sewage Discharges in Unsewered Areas

Livestock (Grazing or Feeding Operations)

Unrestricted Cattle Access

Rural (Residential Areas)

New River Basin

Cause Group Code: N05R-01-BEN Elk Creek and Turkey Fork

Cause Location: This segment includes the mainstem from the confluence of Comers Rock Branch downstream to Turkey Fork. It

also includes the headwaters of Turkey Fork near Dry Run Gap on Iron Mountain.

City / County: Grayson Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 5A

The Probabilistic Monitoring station located at 9-EKC013.81 was impaired based on VSCI scores of 47 and 45. Probabilistic Monitoring station 9-TKY001.55 was impaired based on VSCI scores of 40 and 39.

Assessment Unit / Water Name / Location Desc.	Caus Categ	se ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N05R_EKC03A02 / Elk Creek / Mainstem from confluence Comers Rock Branch near Bennington Mill, downstream to Turkey Fork confluence, WQS Section 2 (NE13).		Benthic Macroinvertebr Bioassessments	ates	2008	L	9.38
VAS-N05R_TKY01A02 / Turkey Fork / Headwaters near Dry Ru Gap on Iron Mountain in Jefferson National Forest, WQS Section (NE14).		Benthic Macroinvertebr Bioassessments	ates	2018	L	6.00
Elk Creek and Turkey Fork			Estuary		servoir	River
Aquatic Life Ronthic Macroinvertebrates Ricassessments Tata	l Impoire	d Siza by Water Type:	(Sq. Miles)	(A	cres)	(Miles) 15.38
Benthic Macroinvertebrates Bioassessments - Total	i iiiipaiie	tu Size by Water Type.				13.30

Sources:

Animal Feeding Operations Gra
(NPS) Graph

Grazing in Riparian or Shoreline Zones

Unrestricted Cattle Access

New River Basin

Cause Group Code: N06R-01-BAC Chestnut Creek

Cause Location: This segment extends from the confluence with Coal Creek downstream to river mile 14.27 at the Galax raw water intake and from river mile 14.27 downstream to the Allied-Signal Gossan mine discharge at river mile 8.06. It also

includes Lower Chestnut Creek from the Skunk Branch confluence at the Allied Gossan mine, river mile 8.06,

downstream to the confluence with New River.

City / County: Carroll Co. Galax City Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The AWQM station, 9-CST0016.82, had a 20% exceedance of the E.coli water quality standard, station 9-CST002.64 25% exceeded WQS. Station 9-CST012.75 had a 50% exceedance of the E.coli water quality standard.

Escherichia coli (E. coli) - Total Im	paired	Size by Water Type:				17.55
Chestnut Creek Recreation			Estuary (Sq. Miles)		servoir cres)	River (Miles)
VAS-N06R_CST04A98 / Chestnut Creek / This is an upstream continuation of the public water supply segment for the City of Galax raw water intake extending upstream to Cox Mill, WQS Section 2h (NE17)	4A	Escherichia coli (E. coli)		2004	L	2.10
VAS-N06R_CST03A94 / Chestnut Creek / Segment extends from the southern Route 89 bridge, river mile 15.00, near the upstream Galax City limit, downstream to river mile 14.27, the Galax raw water ntake, WQS Section 2h (NE17)	4A	Escherichia coli (E. coli)		2004	L	1.09
VAS-N06R_CST02A94 / Chestnut Creek / Segment extends from the City of Galax Water Treatment Plant intake, river mile 14.27, downstream to the Allied-Signal Gossan mine discharge, river mile 3.06, Section 2 (NE17).	4A	Escherichia coli (E. coli)		2016	L	5.68
VAS-N06R_CST01A94 / Chestnut Creek / Lower Chestnut Creek from Skunk Branch confluence at Allied Gossan mine, river mile 3.06, downstream to the confluence with New River, WQS Section 2 (NE17).	4A	Escherichia coli (E. coli)		2014	L	8.68
	Cause atego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size

Sources:

Animal Feeding Operations Crop Production (Crop Land Livestock (Grazing or NPS) Rural (Residential Areas) Feeding Operations)

Source Unknown

New River Basin

Cause Group Code: N06R-01-BEN Chestnut Creek

Cause Location: This segment includes the mainstem of Chestnut Creek from the Skunk Branch confluence downstream to the

confluence with New River.

City / County: Carroll Co. Galax City Grayson Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

The AWQM station, 9-CST002.64, historically indicated an impairment of the aquatic life use. Station 9-CST005.73 has VSCI scores of 49.6 and 65.9 in 2014 and station 9-CST012.63 had VSCI scores of 56.9 and 50.8 in 2013.

_	Cause Catego	e rry Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N06R_CST01A94 / Chestnut Creek / Lower Chestnut Creek from Skunk Branch confluence at Allied Gossan mine, river mile 8.06, downstream to the confluence with New River, WQS Section 2 (NE17).	4A	Benthic Macroinvertebrates Bioassessments	2002	L	8.68
VAS-N06R_CST02A94 / Chestnut Creek / Segment extends from the City of Galax Water Treatment Plant intake, river mile 14.27, downstream to the Allied-Signal Gossan mine discharge, river mile 8.06, Section 2 (NE17).	4A	Benthic Macroinvertebrates Bioassessments	2004	L	5.68

Chestnut Creek	Estuary	Reservoir	River
Aquatic Life	(Sq. Miles)	(Acres)	(Miles)
Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type:			14.36

Sources:

Acid Mine Drainage Crop Production (Crop Land Grazing in Riparian or Mine Tailings

or Dry Land) Shoreline Zones

Silviculture Activities Unrestricted Cattle Access Urban Runoff/Storm Sewers

New River Basin

Cause Group Code: N06R-03-BAC **Meadow Creek & New River**

Cause Location: This segment includes Meadow Creek and its tributaries and New River from Elk Creek confluence downstream to

Eagle Bottom Creek confluence.

City / County: Grayson Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

AWQM station located at 9-MCR000.20 had a 75% exceedance of the E. coli water quality standard. 9-NEW148.23 has E.coli exceedance rate of 23%.

	Cause		Cycle First	TMDL Dev.	Water
Assessment Unit / Water Name / Location Desc.	Category Cause Name		isted	Priority	Size
VAS-N06R_MCR01A02 / Meadow Creek & tributaries / Meadow Creek from confluence with New River upstream to headwaters and tributaries, south west of Galax WQS Section 2, v, NEW-5 (NE15).	5A Escherichia coli (E. coli)	:	2010	M	10.53
VAS-N06R_NEW01A00 / New River / Mainstem from the Elk Creek confluence near Riverside to five miles above Fries Dam, WQS Section 2 (NE15/16).	5A Escherichia coli (E. coli)	1	2004	M	5.38
VAS-N06R_NEW02A02 / New River / New River mainstem from Fries Dam, five miles upstream, Section 2i (NE16).	5A Escherichia coli (E. coli)	;	2014	M	5.03
Meadow Creek & New River		Estuary	Res	ervoir	River
Recreation		(Sq. Miles)	(Ad	cres)	(Miles)
Escherichia coli (E. coli) - Total Ir	npaired Size by Water Type:				20.94

Sources:

Livestock (Grazing or	Rural (Residential Areas)	Source Unknown	Unrestricted Cattle Access
Feeding Operations)			

New River Basin

Cause Group Code: N06R-03-PH New River

Cause Location: Mainstem from the Elk Creek confluence near Riverside to five miles above the Fries Dam and from the Fries

Dame, five miles upstream.

City / County: Grayson Co.

Use(s): Aquatic Life

Cause(s) / VA Category: pH / 5A

18% pf the pH measurements at station 9-NEW148.23 exceed the maximum water quality standard.

pH - Total	Impaired Size by Water Type:				10.41
Aquatic Life		(Sq. Miles)	(Ad	cres)	(Miles)
New River		Estuary	Res	ervoir	River
VAS-N06R_NEW02A02 / New River / New River mainstem from Fries Dam, five miles upstream, Section 2i (NE16).	n 5A pH	2	2018	L	5.03
VAS-N06R_NEW01A00 / New River / Mainstem from the Elk Creek confluence near Riverside to five miles above Fries Dam, WQS Section 2 (NE15/16).	5A pH	2	2018	L	5.38
Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	Ī	Cycle First isted	TMDL Dev. Priority	Water Size

Sources:

Source Unknown

New River Basin

Cause Group Code: N07R-01-BAC Crooked Creek

Cause Location: This segment extends from the headwaters of Crooked Creek downstream to the confluence with New River at

Byllesby.

City / County: Carroll Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

Fecal Coliform / 5A

The AWQM station, 9-CRK020.79, had a 25% exceedance of the E. coli water quality standard. This segment is designated natural trout waters. Station 9-CRK015.69 had a 50% exceedance of the E.coli water quality standard. 9-CRK003.00 has 33% exceedance.

Assessment Unit / Water Name / Location Desc.	Cause Catego	e rry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N07R_CRK01A04 / Crooked Creek / From headwaters near Pipers Gap to Beaverdam Creek confluence south of Woodlawn, WQS Section 2, ii (NE19).	5A	Escherichia coli (E. coli)		2010	М	11.45
VAS-N07R_CRK01A98 / Crooked Creek / From confluence of Cranberry Creek east of SR 635, downstream to New River at Byllesby, WQS Section 2, iii (NE20).	5A	Escherichia coli (E. coli)		2010	M	12.09
VAS-N07R_CRK02A04 / Crooked Creek / From Beaverdam Cree confluence, south of Woodlawn, to Cranberry Creek confluence, WQS Section 2 (NE20).	k 5A	Escherichia coli (E. coli)		2010	M	4.36
Crooked Creek Recreation			Estuary (Sq. Miles		servoir .cres)	River (Miles)
Escherichia coli (E. coli) - Total Ir	npaire	d Size by Water Type:				27.90
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N07R_CRK01A04 / Crooked Creek / From headwaters near Pipers Gap to Beaverdam Creek confluence south of Woodlawn, WQS Section 2, ii (NE19).	5A	Fecal Coliform		2004	М	11.45
VAS-N07R_CRK01A98 / Crooked Creek / From confluence of Cranberry Creek east of SR 635, downstream to New River at Byllesby, WQS Section 2, iii (NE20).	5A	Fecal Coliform		2004	M	12.09
Crooked Creek Recreation			Estuary (Sq. Miles		servoir .cres)	River (Miles)
Fecal Coliform - Total Ir	npaired	d Size by Water Type:	(3400	, (*)	,	23.54

Sources:

Grazing in Riparian or Shoreline Zones

Source Unknown

Unrestricted Cattle Access

New River Basin

Cause Group Code: N07R-01-TEMP **Crooked Creek**

Cause Location: This segment of Crooked Creek begins at Route 707 and continues to Route 620.

City / County: Carroll Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5A

The AWQM station located at 9-CRK015.69 had a 25% exceedance of the temperature standard for natural trout streams.

Cycle **TMDL** First Cause Dev. Water Listed Priority Category Cause Name Size Assessment Unit / Water Name / Location Desc. VAS-N07R_CRK02A04 / Crooked Creek / From Beaverdam Creek 5A Temperature 2010 4.36 confluence, south of Woodlawn, to Cranberry Creek confluence,

WQS Section 2 (NE20).

Crooked Creek **Estuary** Reservoir River (Sq. Miles) (Acres) (Miles) **Aquatic Life**

Temperature - Total Impaired Size by Water Type: 4.36

Sources:

Grazing in Riparian or Shoreline Zones

Unrestricted Cattle Access

New River Basin

Cause Group Code: N08R-01-BAC New River

Cause Location: This segment includes the tributaries of the New River from the Reed Creek confluence downstream to the Big Reed Island Creek confluence including Pine Run. This bacteria impaired section of the New River mainstem is

between the Big Reed Island Creek confluence, near Route 100, and the backwaters of Claytor Lake near the

Fecal Coliform / 5A

Wythe/Pulaski county line.

City / County: Carroll Co. Pulaski Co. Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

Station 9-NEW107.51 was formally contained in the 1999 Federal Consent Decree Attachment B List for fecal coliform bacteria. The 2002 Assessment found the recreational use fully supported. Station 107.51 found only three of 52 samples in excess of the former WQS fecal coliform 1000 n/100 ml instantaneous criterion. The waters were therefore not 303(d) listed in 2002. The initial 303(d) Listing for fecal coliform bacteria occurs with the 2004 IR. The 2006 Assessment and 303(d) Listings replace fecal coliform bacteria with escherichia coli (E.coli) bacteria as the indicator with sufficient E.coli data as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters]. The waters were delisted with the 2014 assessment as E.coli excursions of the 235 cfu/100 ml instantaneous criterion are three of 35 observations. An 8.5% exceedance rate at station 9-NEW107.51 (Allisonia USGS Gage) results in the delisting of these 1999 Consent Decree waters. These waters are listed in the 2016 data window based on the information detailed below.

9-NEW107.51 (Allisonia USGS Gage) The 2016 and 2018 Integrated Reports find five of 39 E.coli samples in excess of the WQS instantaneous criterion of 235 cfu/100 ml. Excessive values range from 280 cfu/100 ml to greater than 2000.

9-PRN000.84 had a 41% (5/12) exceedance of the E. coli water quality standard.

9-MRN000.31 - E.Coli bacteria WQS exceedance rate was 58%.

Assessment Unit / Water Name / Location Desc.	Cause Categor	ry Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N08R_MRN01A04 / Mill Creek / Enters New River from rupper end is near SR 606 near New Jersey Zinc, WQS Section (NE25).		Escherichia coli (E. coli)	2010	M	4.37
VAS-N08R_NEW01A02 / New River / Mainstem, north of Barr Springs, from Reed Creek confluence downstream to Big Reed Island Creek confluence, WQS Section 2 (NE43).	ren 5A	Escherichia coli (E. coli)	2016	M	5.71
VAS-N08R_NEW01B98 / New River / From Mill Creek confluence and Austinville, downstream to the confluence of unnamed tributest of Flatwood, WQS Section 2 (NE21).		Escherichia coli (E. coli)	2016	M	1.44
VAS-N08R_NEW01L98 / New River at Byllesby / New River mainstem in Carroll County. This is a run-of-River power general facility with limited public access that extends from Buck Dam upstream to Byllesby Dam (NE21).		Escherichia coli (E. coli)	2008	M	3.06
VAS-N08R_NEW02B00 / New River / Mainstem public water supply segment for Austinville from Buck Dam tailwaters downst to the Mill Creek confluence, WQS Section 2I (NE21).	5A ream	Escherichia coli (E. coli)	2016	M	5.01
VAS-N08R_NEW03B98 / New River / From Buck Dam, to tailwaters, five miles upstream of Austinville raw water intake, se 2 (NE21).	5A ection	Escherichia coli (E. coli)	2016	M	0.92
VAS-N08R_PNR01A10 / Pine Run / At the Wythe/Pulaski Couline, New River tributary from Pine Run Church downstream, WC Section 2 (NE32).	,	Escherichia coli (E. coli)	2010	M	1.43
VAW-N16R_NEW01A00 / New River / This section of the New River extends from the mouth of Big Reed Island Creek downstruction.		Escherichia coli (E. coli)	2006	М	0.61

New River Basin

Wastes from Pets

to the backwaters of Claytor Lake Class IV sec. 2c (NE43).

New River Recreation			Estuary Reservoir (Sq. Miles) (Acres)		River (Miles)	
	Escherichia coli (E. coli) - T	otal Impaired Size by Water Type:				22.5
Assessment Unit / Water	Name / Location Desc.	Cause Category Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
<u>—</u>	Creek / Enters New River from New Jersey Zinc, WQS Section	•		2004	M	4.37
New River			Estuary		servoir	River
Recreation			(Sq. Miles)	(A	cres)	(Miles)
	Fecal Coliform - T	otal Impaired Size by Water Type:				4.37
ources:						
Grazing in Riparian or	Livestock (Grazing or	Municipal (Urbanized High	Source Unknown			

Wet Weather Discharges (Non-Point Source)

Wildlife Other than

Waterfowl

New River Basin

Cause Group Code: N08R-03-BAC **Shorts Creek and Unnamed Tributary**

Cause Location: This segment includes Shorts Creek and continues until it enters New River at Jackson Ferry. This segment also

includes an unnamed tributary to Shorts Creek that enters at Jackson Ferry and flows west from Rackettown.

City / County: Carroll Co. Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A Fecal Coliform / 5A

The AWQM station, 9-SRT000.12, had a 62% exceedance of the E. coli water quality standard.

Assessment Unit / Water Name / Location Desc.	Cause Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N08R_SRT01A04 / Shorts Creek / Headwaters, south of Poplar Camp Mountain, WQS Section 2 (NE 25).	5A	Escherichia coli (E. coli)		2012	М	3.31
VAS-N08R_SRT01B04 / Shorts Creek / The lower reach of Short Creek, enters New River at Jackson Ferry, WQS Section 2, vi (NE2		Escherichia coli (E. coli)		2010	М	7.07
VAS-N08R_XEE01A06 / Shorts Creek unnamed tributary / Flows west from Rackettown and enters Shorts Creek above Jackson Ferry, WQS Section 2 (NE25).	5 5A	Escherichia coli (E. coli)		2010	M	3.88
Shorts Creek and Unnamed Tributary Recreation			Estuary (Sq. Miles)		servoir cres)	River (Miles)
Escherichia coli (E. coli) - Total	mpaire	d Size by Water Type:				14.26
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N08R_SRT01B04 / Shorts Creek / The lower reach of Short Creek, enters New River at Jackson Ferry, WQS Section 2, vi (NE2		Fecal Coliform		2004	М	7.07
VAS-N08R_XEE01A06 / Shorts Creek unnamed tributary / Flows west from Rackettown and enters Shorts Creek above Jackson Ferry, WQS Section 2 (NE25).	5A	Fecal Coliform		2006	M	3.88
Shorts Creek and Unnamed Tributary			Estuary	Res	servoir	River
Recreation			(Sq. Miles)	(A	cres)	(Miles)
Fecal Coliform - Total I	mpaire	d Size by Water Type:				10.95
Sources:						

Animal Feeding Operations Grazing in Riparian or Livestock (Grazing or **Unrestricted Cattle Access** Shoreline Zones Feeding Operations) (NPS)

New River Basin

Cause Group Code: N09R-01-BAC Cripple Creek and Crigger Creek

Cause Location: This segment includes the mainstem from the confluence with Dry Run, downstream to the Francis Mill Creek confluence as well as the lower segment of the mainstem from the New River confluence upstream to the Dean Branch confluence. It also includes Crigger Creek from the confluence with Cripple Creek upstream to the

confluence with Middle Creek.

City / County: Smyth Co. Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A Fecal Coliform / 4A

The AWQM stations, 9-CPL018.47 and 9-CPL022.99, both had a 45% exceedance of the E. coli water quality standard. AWQM stations 9-CPL0001.03 had a 25% exceedance, 9-CPL003.10 had a 17% exceedance and 9-CGG000.35 had a 36% exceedance of the E. coli water quality standard.

	Cause			Cycle First	TMDL Dev.	Water
Assessment Unit / Water Name / Location Desc.	Catego	ry Cause Name		Listed	Priority	Size
$VAS-N09R_CGG01B04\ /\ Crigger\ Creek\ /\ From\ confluence\ with\ Cripple\ Creek\ upstream\ to\ Middle\ Creek\ confluence,\ WQS\ Section\ iv\ (NE22).$	4A 2,	Escherichia coli (E. coli)		2018	L	4.20
VAS-N09R_CPL01A98 / Cripple Creek / Extends from Dean Branch confluence upstream to Francis Mill Creek confluence, WQS Section 2 (NE24).	4A S	Escherichia coli (E. coli)		2018	L	11.68
VAS-N09R_CPL01B04 / Cripple Creek / Lower segment of mainstem from the New River confluence upstream to the Dean Branch confluence at Porter Crossroads, WQS Section 2I (NE24).	4A	Escherichia coli (E. coli)		2010	L	3.17
VAS-N09R_CPL02A98 / Cripple Creek / From the Dry Run confluence near Speedwell downstream to the Francis Mill Creek confluence, WQS Section 2 (NE23).	4A	Escherichia coli (E. coli)		2010	L	6.49
VAS-N09R_CPL02B04 / Cripple Creek / Mainstem from Blue Spring Creek confluence downstream to the Dry Run confluence near Speedwell, WQS Section 2, *** (NE23).	4A	Escherichia coli (E. coli)		2010	L	6.43
Cripple Creek and Crigger Creek			Estuary	Res	servoir	River
Recreation			(Sq. Miles)	(A	cres)	(Miles)
Escherichia coli (E. coli) - Total I	mpaired	Size by Water Type:				31.97
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N09R_CPL01B04 / Cripple Creek / Lower segment of mainstem from the New River confluence upstream to the Dean Branch confluence at Porter Crossroads, WQS Section 2I (NE24).	4A	Fecal Coliform		2004	L	3.17
VAS-N09R_CPL02A98 / Cripple Creek / From the Dry Run confluence near Speedwell downstream to the Francis Mill Creek confluence, WQS Section 2 (NE23).	4A	Fecal Coliform		2004	L	6.49
Cripple Creek and Crigger Creek			Estuary		servoir	River
Recreation			(Sq. Miles)	(A	cres)	(Miles)
Fecal Coliform - Total I	Impaired	Size by Water Type:				9.66

New River Basin

Sources:

Livestock (Grazing or Feeding Operations)

Unrestricted Cattle Access

New River Basin

Cause Group Code: N09R-03-BAC Slate Spring Branch and Dean Branch

Cause Location: This segment includes Slate Spring Branch from the Cripple Creek confluence up stream to the headwaters and

Dean Branch from the confluence with Cripple Creek upstream 1.7 miles.

City / County: Smyth Co. Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A Fecal Coliform / 4A

The AWQM station, 9-SPB000.10, had a 100% exceedance of the E.coli water quality standard. Station 9-DEN000.03 had a

25% exceedance of the E.coli water quality standard.

Cause Assessment Unit / Water Name / Location Desc. Cause Category Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N09R_DEN01A10 / Dean Branch / Cripple Creek tributary at 4A Escherichia coli (E. coli) Porters Crossroads, WQS Section 2 (NE24).		2010	L	1.92
VAS-N09R_SPB01A04 / Slate Spring Branch / From Cripple Creek 4A Escherichia coli (E. coli) confluence at Eagle Cliff upstream to headwaters at Matney Flat, WQS Section 2.		2010	L	6.14
Slate Spring Branch and Dean Branch	Estuary		servoir	River
Recreation	(Sq. Miles) (A	cres)	(Miles)
Escherichia coli (E. coli) - Total Impaired Size by Water Type:				8.06
Cause Assessment Unit / Water Name / Location Desc. Category Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N09R_SPB01A04 / Slate Spring Branch / From Cripple Creek 4A Fecal Coliform confluence at Eagle Cliff upstream to headwaters at Matney Flat, WQS Section 2.		2004	L	6.14
Slate Spring Branch and Dean Branch Recreation	Estuary (Sq. Miles		servoir cres)	River (Miles)
Fecal Coliform - Total Impaired Size by Water Type:				6.14

Sources:

Animal Feeding Operations Non-Point Source Source Unknown Unrestricted Cattle Access (NPS)

New River Basin

Cause Group Code: N09R-03-BEN **Dean Branch**

Cause Location: A Cripple Creek tributary at Porters Crossroads.

City / County: Smyth Co. Wythe Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 5A

The freshwater probabilistic monitoring station at 9-DEN000.39 was impaired based on VSCI scores of 54.70 and 57.54.

Cycle **TMDL** First Cause Dev. Water Listed Priority Category Cause Name Assessment Unit / Water Name / Location Desc. Size 5A 2016 1.92 VAS-N09R_DEN01A10 / Dean Branch / Cripple Creek tributary at Benthic Macroinvertebrates Bioassessments Porters Crossroads, WQS Section 2 (NE24). Dean Branch Estuary Reservoir River

(Sq. Miles) (Acres) (Miles) **Aquatic Life** 1.92

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type:

Sources:

Non-Point Source **Animal Feeding Operations** Livestock (Grazing or Feeding Operations) (NPS)

New River Basin

Cause Group Code: N09R-04-TEMP Crigger Creek

Cause Location: From the confluence with Cripple Creek upstream to the Middle Creek confluence.

City / County: Smyth Co. Wythe Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5A

Station 9-CGG000.35 had 2 of 12 (17%) temperature measurements exceed the WQS for Class VI waters.

Cause Cycle TMDL
Cause First Dev. Water
Assessment Unit / Water Name / Location Desc. Category Cause Name Listed Priority Size

VAS-N09R_CGG01B04 / Crigger Creek / From confluence with 5A Temperature 2020 L 4.20

Cripple Creek upstream to Middle Creek confluence, WQS Section 2,

iv (NE22).

Crigger CreekEstuaryReservoirRiverAquatic Life(Sq. Miles)(Acres)(Miles)

Temperature - Total Impaired Size by Water Type: 4.20

Sources:

Grazing in Riparian or Source Unknown Unrestricted Cattle Access

Shoreline Zones

New River Basin

Cause Group Code: N10R-01-TEMP Reed Creek and Mill Creek

Cause Location: Reed Creek mainstem from Venrick Run upstream to South Fork and Mill Creek from its headwaters west of Rural

Retreat to the Reed Creek confluence east of Blacklick.

City / County: Wythe Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5A

Water temperature was exceeded for Class VI WQS at 9-RDC033.94 in 2 of 9 measurements. Station 9-MCE000.37 had 2 of

Cycle TMDI

12 temperature measurements exceed the WQS for Class VI waters.

Temperature - Total	Impaired Size by Water Type:				7.82
Reed Creek and Mill Creek Aquatic Life		Estuary (Sq. Miles)		ervoir cres)	River (Miles)
VAS-N10R_RDC01A00 / Reed Creek / Reed Creek mainstem parallel to SR 659 from Venrick Run upstream to South Fork confluence south of Petunia in Section 2g (NE28).	5A Temperature	2	012	L	1.43
VAS-N10R_MCE01A02 / Mill Creek / From headwaters west of Rural Retreat to Reed Creek confluence east of Blacklick, WQS Section 2.	5A Temperature	2	020	L	6.39
Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	F	irst sted	Dev. Priority	Water Size

Sources:

Grazing in Riparian or Shoreline Zones

Loss of Riparian Habitat

New River Basin

Cause Group Code: N10R-02-BAC South Fork Reed Creek and Mill Creek

Cause Location: This segment includes the mainstem of South Fork Reed Creek downstream to the Reed Creek confluence as well as the mainstem of Mill Creek to the confluence with Reed Creek. It also includes Hubble Branch, north of I81 near

Rural Retreat.

City / County: Smyth Co. Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The AWQM station, 9-MCE000.37, had a 66% exceedance of the E.coli water quality standard. Station 9- RSF000.08 & 9-RSF006.78 had a 67% exceedance of the E.coli water quality standard. 9-HOL000.74 had 66% E.coli exceedance rate.

Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	Cyc Firs List	st Dev.	Water Size
VAS-N10R_HOL01A12 / Huddle Branch / A Mill Creek tributary from the Monkey Run confluence parallel SR 617 North of I81 at Staley Crossroads (NE26).	4A Escherichia coli (E. coli) 201	2 L	1.48
VAS-N10R_MCE01A02 / Mill Creek / From headwaters west of Rural Retreat to Reed Creek confluence east of Blacklick, WQS Section 2.	4A Escherichia coli (E. coli) 200	16 L	6.39
VAS-N10R_RSF01A00 / South Fork Reed Creek / Mainstem from river mile 6.8 near Groseclose, downstream to the Reed Creek confluence parallel and south of I81, WQS Section 2; Wytheville National Fish Hatchery is on this reach (NE28).	m 4A Escherichia coli (E. coli) 200	06 L	6.77
VAS-N10R_RSF01A02 / South Fork Reed Creek / Mainstem in headwaters near Fairview and through Groseclose, WQS Section 2 vi (NE28).	4A Escherichia coli (E. coli 2,) 201	2 L	13.35
South Fork Reed Creek and Mill Creek Recreation		Estuary (Sq. Miles)	Reservoir (Acres)	River (Miles)
Escherichia coli (E. coli) - Total I	Impaired Size by Water Type:	· · · /	(* ::::55)	27.99

Sources:

Animal Feeding Operations	Grazing in Riparian or	Livestock (Grazing or	Unrestricted Cattle Access
(NPS)	Shoreline Zones	Feeding Operations)	

New River Basin

Cause Group Code: N10R-02-BEN Mill Creek

Cause Location: From the headwaters, west of Rural Retreat, to the confluence with Reed Creek, east of Blacklick.

City / County: Wythe Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 5A

The probabilistic monitoring station at 9-MCE000.27 was impaired based on VSCI scores of 58 and 51.

Assessment Unit / Water Name / Location Desc.	Cause	First	Dev.	Water
	Category Cause Name	Listed	Priority	Size
VAS-N10R_MCE01A02 / Mill Creek / From headwaters west of Rural Retreat to Reed Creek confluence east of Blacklick, WQS Section 2.	5A Benthic Macroinvertebrates Bioassessments	2018	L	6.39

Mill Creek Estuary Reservoir River (Sq. Miles) (Acres) (Miles) **Aquatic Life** 6.39

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type:

Sources:

Source Unknown

New River Basin

Cause Group Code: N10R-03-BAC Stony Fork

Cause Location: This segment includes the headwaters downstream to the Reed Creek confluence.

City / County: Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The AWQM station, 9-SFK000.28, had a 67% exceedance of the E. coli water quality standard and station 9-SFK001.51 had a 50% exceedance of the E.coli water quality standard.

Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	Cycle First Liste	t Dev.	Water Size
VAS-N10R_SFK01A02 / Stony Fork / Class V waters @ Favonia downstream to Reed Creek confluence, WQS Section 2, vi (NE27)	•) 2006	L	1.90
VAS-N10R_SFK01A12 / Stony Fork / Headwaters in Jefferson National Forest south of Walker Mountain downstream to Class VI waters @ Favonia, WQS Section 2, vi (NE27).	4A Escherichia coli (E. coli) 2012	. L	4.73
Stony Fork			Reservoir	River
Pecreation		(Sa. Miles)	(Acres)	(Miles)

Stony Fork	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Escherichia coli (E. coli) - Tota	I Impaired Size by Water Type:		6.63

Sources:

Livestock (Grazing or Feeding Operations)

Unrestricted Cattle Access

New River Basin

Cause Group Code: N10R-04-BAC Tate Run

Cause Location: This segment begins at the Stuffle Run confluence and extends downstream to Reed Creek.

City / County: Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The AWQM station, 9-TAT000.46, had a 58% exceedance of the E. coli water quality standard.

Cause Cycle TMDL
First Dev. Water
Assessment Unit / Water Name / Location Desc. Category Cause Name Listed Priority Size

VAS-N10R_TAT01A06 / Tate Run / From Stuffle Run confluence 4A Escherichia coli (E. coli) 2006 L 0.56

downstream to Reed Creek, Section 2g (NE28).

Tate Run Estuary Reservoir River (Sq. Miles) (Acres) (Miles)

Escherichia coli (E. coli) - Total Impaired Size by Water Type: 0.56

Sources:

Livestock (Grazing or Unrestricted Cattle Access Feeding Operations)

New River Basin

Cause Group Code: N11R-01-BAC Reed Creek and Pine Run

Cause Location: This segment begins at the Gullion Fork confluence and extends downstream to the Venrick Run confluence. It also includes the lower mainstem of Reed Creek from its confluence with an unnamed tributary East of Route 21 to the confluence with Miller Creek and from the New River confluence near Lone Ash, upstream to the Glade Creek

confluence. Pine Run, a Reed Creek tributary north of I-81 and south of Pine Ridge.

City / County: Smyth Co. Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

Fecal Coliform / 4A

The AWQM station, 9-RDC046.65, had a 33% exceedance of the E.coli water quality standard. Station 9-RDC038.01 had a 25% exceedance of the E. coli standard and station 9-RDC033.78 had a 50% exceedance and station 9-RDC023.24 had a 27% exceedance of the E.coli water quality standard. Station 9-RDC009.00 had a 14% exceedance of the e.coli WQS and station 9-PRN000.04 had a 67% exceedance.

Assessment Unit / Water Name / Location Desc. Co /AS-N10R_RDC01A00 / Reed Creek / Reed Creek mainstem parallel to SR 659 from Venrick Run upstream to South Fork	atego 4A	ry Cause Name Fecal Coliform		Listed 2002	Priority L	Size 1.43
	Cause			Cycle First	TMDL Dev.	Water
Escherichia coli (E. coli) - Total Imp	paired	Size by Water Type:				49.40
Reed Creek and Pine Run Recreation			Estuary (Sq. Miles)		servoir .cres)	River (Miles)
/AS-N11R_RDC03B04 / Reed Creek / From New River confluence near Lone Ash, upstream to the Glade Creek confluence near Boiling Spring, WQS Section 2 (NE31).	4A	Escherichia coli (E. coli)		2004	L	9.87
/AS-N11R_RDC01C02 / Reed Creek / Segment begins at confluence of unnamed tributary east of Rt. 21 bridge and extends downstream to the Muskrat Branch confluence, north of Rt. 11, WQS Section 2 (NE29).	4A	Escherichia coli (E. coli)		2010	L	6.21
/AS-N11R_RDC01B00 / Reed Creek / Lower mainstem from Muskrat Branch confluence downstream to Rt. 52 bridge south of Max Meadows, WQS Section 2 (NE29).	4A	Escherichia coli (E. coli)		2006	L	5.85
/AS-N10R_RDC01C02 / Reed Creek / Headwaters of Reed Creek rom Redding Gap in Jefferson National Forest downstream to Gullion Fork confluence, WQS Section 2, *** (NE26).	4A	Escherichia coli (E. coli)		2012	L	6.83
/AS-N10R_RDC01B00 / Reed Creek / Mainstem from the Stony Fork confluence south of Favonia, upstream to the Gullion Fork confluence, WQS Section 2, *** (NE26).	4A	Escherichia coli (E. coli)		2006	L	9.85
/AS-N10R_RDC01A02 / Reed Creek / From South Fork Reed Creek confluence upstream to Stony Fork confluence west of Petunia, WQS Section 2, *** (NE28).	4A	Escherichia coli (E. coli)		2006	L	5.23
/AS-N10R_RDC01A00 / Reed Creek / Reed Creek mainstem parallel to SR 659 from Venrick Run upstream to South Fork confluence south of Petunia in Section 2g (NE28).	4A	Escherichia coli (E. coli)		2008	L	1.43
/AS-N10R_PRN01A12 / Pine Run / Reed Creek tributary north of 81 and south of Pine Ridge (NE28).	4A	Escherichia coli (E. coli)		2012	L	4.13
	Cause atego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size

Final 2020 Appendix 5 - 3416

confluence south of Petunia in Section 2g (NE28).

New River Basin

Reed Creek and Pine Run

Estuary
Recreation

Reservoir
River
(Sq. Miles)
(Acres)
(Miles)

Fecal Coliform - Total Impaired Size by Water Type:

1.43

Sources:

Animal Feeding Operations (NPS)

Unrestricted Cattle Access

Grazing in Riparian or Shoreline Zones

Livestock (Grazing or Feeding Operations)

Source Unknown

New River Basin

Cause Group Code: N11R-02-BAC Miller Creek

Cause Location: This segment includes the mainstem from the Beaverdam confluence at Max Meadows downstream to Reed Creek

and from the West Fork confluence downstream to Max Meadows.

City / County: Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The AWQM station, 9-MER000.09, had a 45% exceedance and station 9-MER000.85 had a 20% of the E.coli water quality

standard.

Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size	
VAS-N11R_MER01A06 / Miller Creek / From Beaverdam confluence in the community of Max Meadows downstream to Ree Creek, WQS Section 2 (NE31).	4A d	Escherichia coli (E. coli)		2006	L	0.42	
VAS-N11R_MER02A10 / Miller Creek / A Reed Creek tributary From West Fork confluence on Brushy Ridge downstream to Max Meadows, WQS Section 2 (NE31).	4A	Escherichia coli (E. coli)		2012	L	3.64	
Miller Creek			Estuary	Res	servoir	River	
Recreation			(Sq. Miles)	(A	cres)	(Miles)	
Escherichia coli (E. coli) - Total Impaired Size by Water Type:							

Sources:

Rural (Residential Areas) Unrestricted Cattle Access

New River Basin

Cause Group Code: N11R-02-BEN Reed Creek tributary

Cause Location: This segment includes an unnamed tributary of Reed Creek that drains the Wytheville Community College at the

east end of the town of Wytheville.

City / County: Wythe Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 5A

Station 9-XES000.94 was impaired based on VASCI scores of 41 and 51 in 2008.

TMDL Cycle First Dev. Cause Water Listed **Priority** Size Assessment Unit / Water Name / Location Desc. Category Cause Name VAS-N11R_XES01A10 / Reed Creek tributaries / Tributary that 5A Benthic Macroinvertebrates 2010 Μ 2.67 Bioassessments

drains location of Wytheville Community College at east end of Wytheville, WQS Section 2 (NE29).

Reed Creek tributary **Estuary** Reservoir River (Sq. Miles) (Acres) (Miles) **Aquatic Life**

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type:

2.67

Sources:

Grazing in Riparian or Rural (Residential Areas) Streambank Urban Runoff/Storm Sewers Shoreline Zones Modifications/Destabilization

New River Basin

Cause Group Code: N11R-03-BAC McGavock Creek

Cause Location: A Reed Creek tributary east of Grahams Forge, parallel to Route 618.

City / County: Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The AWQM Station located at 9-MGV000.37 has a 18% exceedance of the E. coli water quality standard.

Cycle **TMDL** First Cause Dev. Water Priority Listed Size Assessment Unit / Water Name / Location Desc. Category Cause Name 2012 2.58 VAS-N11R_MGV01A12 / McGavock Creek / Reed Creek tributary 4A Escherichia coli (E. coli) west of Grahams Forge and parallel, to SR 618 (NE31).

McGavock Creek Estuary Reservoir River (Sq. Miles) (Acres) (Miles) Recreation 2.58

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

Sources:

Grazing in Riparian or Livestock (Grazing or Shoreline Zones Feeding Operations)

New River Basin

Cause Group Code: N12R-01-BAC Cove Creek and St. Lukes Fork

Cause Location: This segment includes the lower Cove Creek mainstem from St. Lukes Fork downstream to the confluence with

Reed Creek. This segment also includes St. Lukes Fork from the Cove Creek confluence upstream 1.4 miles, north

of Queens Knob.

City / County: Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The AWQM station, 9-CVR003.88, had a 45% exceedance of the E.coli water quality standard. Station 9-SLK001.24 had a 82% exceedance of the E.coli water quality standard.

Escherichia coli (E. coli) - Total Imp	paired	Size by Water Type:				11.69
Cove Creek and St. Lukes Fork Recreation			Estuary (Sq. Miles)		servoir .cres)	River (Miles)
VAS-N12R_SLK01A04 / St. Lukes Fork / From Cove Creek confluence upstream 1.4 miles, north of Queens Knob, in Section 2 (NE30).	4A	Escherichia coli (E. coli)		2016	L	1.77
VAS-N12R_CVR01A00 / Cove Creek / Lower Cove Creek from St. Lukes Fork confluence, near Queens Knob, downstream to the confluence with Reed Creek, east of Wytheville, WQS Section 2 (NE30).	4A	Escherichia coli (E. coli)		2006	L	9.92
	Cause atego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size

Sources:

Livestock (Grazing or Feeding Operations)

Unrestricted Cattle Access

New River Basin

Cause Group Code: N13R-01-BAC **Big Reed Island Creek**

Cause Location: This segment begins at the headwaters of Big Reed Island Creek and continues downstream to the confluence with

Pine Creek.

City / County: Carroll Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The AWQM station located at 9-RIC039.71 had a 33% exceedance and station 9-RIC049.29 had a 50% exceedance of the

E.coli water quality standard.

Cause Assessment Unit / Water Name / Location Desc. Category Cause Name	Cyc Firs Liste	st Dev.	Water
VAS-N13R_RIC01A00 / Big Reed Island Creek / North of Crooked 5A Escherichia coli (E. coli) Oak from Pine Creek confluence to Snake Creek confluence, WQS Section 2, iii (NE36).	201	8 M	6.64
VAS-N13R_RIC01B04 / Big Reed Island Creek / From headwaters 5A Escherichia coli (E. coli) on Hurricane Knob downstream to Pine Creek confluence near Crooked Oak, WQS Section 2, iii (NE36).	200	8 M	19.85
Big Reed Island Creek	Estuary	Reservoir	River

(Sq. Miles) (Acres) (Miles) Recreation Escherichia coli (E. coli) - Total Impaired Size by Water Type: 26.49

Sources:

Animal Feeding Operations Grazing in Riparian or Shoreline Zones (NPS)

New River Basin

Cause Group Code: N13R-01-BEN Big Reed Island Creek

Cause Location: This segment begins at the headwaters of Big Reed Island Creek and continues downstream to the confluence with

Pine Creek.

City / County: Carroll Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 5A

The Probabilistic Monitoring station located at 9-RIC051.80 was impaired based on the VSCI scores of 70 and 46.

TMDL Cycle First Dev. Cause Water Listed **Priority** Size Category Cause Name Assessment Unit / Water Name / Location Desc. VAS-N13R_RIC01B04 / Big Reed Island Creek / From headwaters 5A Benthic Macroinvertebrates 2008 Н 19.85 Bioassessments on Hurricane Knob downstream to Pine Creek confluence near

Crooked Oak, WQS Section 2, iii (NE36).

Big Reed Island Creek

Aquatic Life

Estuary Reservoir River
(Sq. Miles) (Acres) (Miles)

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type:

19.85

Sources:

Animal Feeding Operations Grazing in Riparian or Unrestricted Cattle Access (NPS) Shoreline Zones

New River Basin

Cause Group Code: N13R-01-TEMP Big Reed Island Creek

Cause Location: From the headwaters on Hurricane Knob downstream to the Pine Creek confluence near Crooked Oak and from

North of Crooked Oak from the Pine Creek confluence to the Snake Creek confluence.

City / County: Carroll Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5A

AWQM station located 9-RIC039.71 had a 25% exceedance and station 9-RIC049.29 had a 25% exceedance of the WQS for

Cycle TMDI

Class IV waters.

Temperature - Total II	mpaired Size by Wat	er Type:		26.49
Big Reed Island Creek Aquatic Life		Estuary (Sq. Miles)	Reservoir (Acres)	River (Miles)
VAS-N13R_RIC01B04 / Big Reed Island Creek / From headwater on Hurricane Knob downstream to Pine Creek confluence near Crooked Oak, WQS Section 2, iii (NE36).	rs 5A Temperature	20	20 L	19.85
VAS-N13R_RIC01A00 / Big Reed Island Creek / North of Crooke Oak from Pine Creek confluence to Snake Creek confluence, WQS Section 2, iii (NE36).		20	18 L	6.64
Assessment Unit / Water Name / Location Desc.	Cause Category Cause Na	Fii me Lis	rst Dev.	Water Size

Sources:

Grazing in Riparian or Shoreline Zones

New River Basin

Cause Group Code: N13R-02-BAC **Snake Creek**

Cause Location: From the Big Reed Island confluence upstream 3.5 miles to near the Macey Branch confluence, WQS Section 2, iii.

City / County: Carroll Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The AWQM station located at 9-SKE000.98 had a 73% exceedance of the E.coli water quality standard.

Assessment Unit / Water Name / Location Desc.	Cause	First	Dev.	Water
	Category Cause Name	Listed	Priority	Size
VAS-N13R_SKE01A04 / Snake Creek / From Big Reed Island Creek confluence upstream 3.5 miles to near Macey Branch confluence, WQS Section 2, iii (NE37).	5A Escherichia coli (E. coli)	2014	М	3.54

Snake Creek Estuary Reservoir River (Sq. Miles) (Acres) (Miles) Recreation Escherichia coli (E. coli) - Total Impaired Size by Water Type: 3.54

Sources:

Source Unknown

New River Basin

Cause Group Code: N14R-01-BAC Big Reed Island Creek

Cause Location: This segment includes the mainstem of Big Reed Island Creek from the confluence of Snake Creek downstream to the confluence with Bobbitt Creek, from Bobbitt Creek to the Greasy Creek confluence, and from the Island Creek

confluence downstream to the Big Branch confluence.

City / County: Carroll Co. Pulaski Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A Fecal Coliform / 5A

The AWQM station, 9-RIC029.23, had a 25% exceedance of the fecal coliform water quality standard and station 9-RIC018.90 had a 17% exceedance of the E. coli water quality standard.

Assessment Unit / Water Name / Location Desc.	Cause Catego	ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N14R_RIC01A00 / Big Reed Island Creek / Big Reed Island Creek east of Red Hill, from Bobbitt Creek confluence upstream to Snake Creek confluence, WQS Section 2, iii (NE39).	5A	Escherichia coli (E. coli)		2018	Н	7.55
VAS-N14R_RIC01B04 / Big Reed Island Creek / Big Reed Island Creek from Bobbitt Creek confluence south of Witcher Knob to Greasy Creek confluence, WQS Section 2 (NE39).	5A	Escherichia coli (E. coli)		2010	Н	13.81
Big Reed Island Creek Recreation			Estuary (Sq. Miles)		servoir cres)	River (Miles)
Escherichia coli (E. coli) - Total I	mpaired	Size by Water Type:				21.36
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N14R_RIC01A00 / Big Reed Island Creek / Big Reed Island Creek east of Red Hill, from Bobbitt Creek confluence upstream to Snake Creek confluence, WQS Section 2, iii (NE39).	5A	Fecal Coliform		2004	Н	7.55
VAS-N14R_RIC01B04 / Big Reed Island Creek / Big Reed Island Creek from Bobbitt Creek confluence south of Witcher Knob to Greasy Creek confluence, WQS Section 2 (NE39).	5A	Fecal Coliform		2014	Н	13.81
Big Reed Island Creek			Estuary		servoir	River
Recreation Fecal Coliform - Total II	mpaired	Size by Water Type:	(Sq. Miles)) (A	cres)	(Miles) 21.36
Fecal Colliorm - Total I	прапес	i Size by water Type:				21.36

Sources:

Source Unknown

New River Basin

Cause Group Code: N14R-01-TEMP Big Reed Island Creek

Cause Location: Big Reed Island Creek east of Red Hill, from the Bobbitt Creek confluence upstream to the Snake Creek confluence.

City / County: Carroll Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5A

AWQM station located 9-RIC029.23 had a 25% exceedance of the WQS for Class IV waters.

Cause Cycle TMDL
Cause First Dev. Water
Assessment Unit / Water Name / Location Desc. Category Cause Name Listed Priority Size

VAS-N14R_RIC01A00 / Big Reed Island Creek / Big Reed Island 5A Temperature 2018 L 7.55

Creek east of Red Hill from Robbitt Creek confluence unstream to

Creek east of Red Hill, from Bobbitt Creek confluence upstream to

Snake Creek confluence, WQS Section 2, iii (NE39).

Big Reed Island Creek

Aquatic Life

Estuary Reservoir River
(Sq. Miles) (Acres) (Miles)

Temperature - Total Impaired Size by Water Type: 7.55

Sources:

Loss of Riparian Habitat

New River Basin

Cause Group Code: N14R-02-BAC **Greasy Creek**

Cause Location: This segment begins at the Carroll county line and continues downstream to the confluence with Big Reed Island

City / County: Floyd Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The AWQM station located at 9-GSC000.03 had a 17% exceedance of the E.coli water quality standard.

TMDL Cycle First Dev. Cause Water Listed **Priority** Size Category Cause Name Assessment Unit / Water Name / Location Desc. VAS-N14R_GSC01A08 / Greasy Creek / From Carroll/Floyd 5A Escherichia coli (E. coli) 2008 Н 13.63 County line downstream to Big Reed Island Creek confluence south

of Macks Mountain, WQS Section 2 (NE40).

Greasy Creek Estuary Reservoir River (Sq. Miles) (Acres) (Miles) Recreation 13.63

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

Sources:

Grazing in Riparian or Shoreline Zones

New River Basin

Cause Group Code: N14R-03-BAC Island Creek

Cause Location: A Big Reed Island Creek tributary northeast of Hillsville from its headwaters near Huffman Knob.

City / County: Carroll Co. Floyd Co. Pulaski Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The AWQM station located at 9-ISL003.05 had a 58% exceedance of the E. coli water quality standard.

Cause Cycle TMDL
Cause First Dev. Water
Assessment Unit / Water Name / Location Desc. Category Cause Name Listed Priority Size

VAS-N14R_ISL01A12 / Island Creek & tributaries / Big Reed 5A Escherichia coli (E. coli) 2018 L 13.35

Island Creek tributary northeast of Hillsville from headwaters near Huffman Knob (NE39).

Island CreekEstuaryReservoirRiverRecreation(Sq. Miles)(Acres)(Miles)

Escherichia coli (E. coli) - Total Impaired Size by Water Type: 13.35

Sources:

Source Unknown

New River Basin

Cause Group Code: N15R-01-BAC Little Reed Island Creek

Cause Location: This segment begins 5 miles above the Hillsville public water intake and extends downstream to the confluence with

Big Reed Island Creek.

City / County: Carroll Co. Pulaski Co. Wythe Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

AWQM station 9-LRI001.62 had a 33% exceedance of the E.coli water quality standard, station 9-LRI009.11 had a 25% exceedance, station 9-LRI017.64 had a 25% exceedance, station 9-LRI023.48 had a 50% exceedance, and station 9-LRI031.58 had a 33% exceedance of the E. coli water quality standard.

Cycle TMDI

Escherichia coli (E. coli) - Total Im	paired	Size by Water Type:				35.98
Little Reed Island Creek Recreation			Estuary (Sq. Miles)		servoir cres)	River (Miles)
/AS-N15R_LRI02A08 / Little Reed Island Creek / Segment extends from Rock Creek confluence upstream to Hillsville PWS intake west of Rt. 100, WQS Section 2 (NE33/34).	5A	Escherichia coli (E. coli)		2008	Н	19.70
/AS-N15R_LRI01B98 / East Fork Little Reed Island Creek / From Hillsville PWS intake south of Rt. 58, upstream five miles, WQS Section 2f (NE33).	5A	Escherichia coli (E. coli)		2008	Н	5.28
/AS-N15R_LRI01A98 / Little Reed Island Creek / Little Reed sland Creek mainstem from confluence with Big Reed Island Creek upstream to Rock C confluence in Carroll County, WQS Section 2 NE34).	5A	Escherichia coli (E. coli)		2008	Н	11.00
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		First Listed	Dev. Priority	Water Size

Sources:

Livestock (Grazing or Feeding Operations)

On-site Treatment Systems
(Septic Systems and Similar Decentralized Systems)

Unspecified Domestic Waste

New River Basin

Cause Group Code: N15R-01-TEMP Little Reed Island Creek

Cause Location: This segment begins approximately 1 mile below the Hillsville water intake and continues downstream to the Big

Reed Island Creek confluence and from the Hillsville PWS intake south of Rt. 58, upstream 5 miles.

City / County: Carroll Co. Pulaski Co. Wythe Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5A

AWQM station 9-LRI017.64 had a 33% exceedance of the temperature standard. Stations 9-LRI020.76 and 9-LRI023.48 had a 25% and 15% exceedance of the temperature standard. Station 9-LRI031.58 had a 25% exceedance of the WQS for Class VI waters

Temperature - Total Impaired Size by Water Type:							
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)			
Little Reed Island Creek		Estuary	Reservoir	River			
VAS-N15R_LRI02A08 / Little Reed Island Creek / Segment extends from Rock Creek confluence upstream to Hillsville PWS intake west of Rt. 100, WQS Section 2 (NE33/34).	5A Temperature	200	8 H	19.70			
VAS-N15R_LRI01B98 / East Fork Little Reed Island Creek / From Hillsville PWS intake south of Rt. 58, upstream five miles, WQS Section 2f (NE33).	5A Temperature	202	0 L	5.28			
Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	Cyc Firs Liste	st Dev.	Water Size			

Sources:

Loss of Riparian Habitat

New River Basin

Cause Group Code: N16L-01-DO Claytor Lake - New River

Cause Location: Claytor Lake - New River mainstem from the mouth of Peak Creek downstream to Claytor Dam (Dublin and Radford

South Quads).

City / County: Pulaski Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Dissolved Oxygen / 4C

A portion of Claytor Lake, 1,799.25 acres, is originally 2002 303(d) Listed for excursions of the Class IV Water Quality Standard (WQS) dissolved oxygen minimum criterion of 4.0 mg/l. The impairment is categorized as natural (4C) in past assessment cycles where no excursions of the Claytor Lake criterion for chlorophyll a (25 μg/L) or total phosphorus (20 μg/L algaecides applied) occur from stations 9-NEW089.34 or 9-NEW087.14 (Lacustrine zone). Virginia's Lake Nutrient Criteria (9 VAC 25-260-187) states the nutrient criteria apply only in the epilimnion for lacustrine waters during thermal stratification for control of nutrient enrichment. Guidance Memo No. 09-2005 "Monitoring and Assessment of Lakes and Reservoirs" outlines criteria for evaluating dissolved oxygen during periods of thermal stratification. Data from the following stations find the waters not supporting the Aquatic Life Use in the epilimnion from dissolved oxygen exceedances of the minimum 4.0 mg/l criterion.

9-NEW092.66- (Dublin Water Works) 2020 epilimnion dissolved oxygen (DO) measurements are 117 exceeding values from a total of 710 measurements. 2018 epilimnion dissolved oxygen (DO) measurements are 113 exceeding values from a total of 635 measurements. 2016 epilimnion dissolved oxygen (DO) measurements are 68 exceeding values from a total of 851 measurements. However these data are not deemed sufficient for delisting these waters. The 2014 data window reports 88 of 787 DO total measurements exceed the minimum 4.0 mg/l criterion. 2012 data reveal 118 of 807 DO measurements exceeding the 4.0 mg/l minimum criterion. The 2010 assessment reports 101 epilimnion dissolved oxygen (DO) measurements exceeding the 4.0 mg/l minimum from 806 measurements. 2008 results find 154 exceed from 656 total observations.

9-NEW089.34- (Line Between Beach and Inlet) 2020 epilimnion DO measurements are 160 of 676 measurements. 2018 epilimnion DO measurements are 161 of 624 measurements. 2016 epilimnion DO measurements are 58 of 806 indicating support of the minimum DO criterion. However these data alone are not sufficient to delist this section of the Lake. Epilimnion DO measurements within the 2014 data window record 59 of 747 total measurements exceed the minimum of 4.0 mg/l. 2012 data exceed in 82 of 798 total measurements. The 2010 assessment finds 99 of 857 epilimnion DO measurements in excess of the minimum criterion. 2008 results find 121 exceed from 637 total observations.

9-NEW087.14- (Under Power Lines above Dam) 2020 epilimnion DO measurements are 155 exceeding of 632 measurements. 2018 epilimnion DO measurements are 164 exceeding of 664 measurements. 2016 epilimnion DO measurements are 58 of 806 indicating support of the minimum DO criterion. However these data alone are not sufficient to delist this section of the Lake. 2012 measurements of DO in the epilimnion are 93 of 804 exceeding the 4.0 minimum criterion. DO exceeds the minimum criterion in 99 of 830 epilimnion measurements within the 2010 data window. 2008 results find 115 exceed from 695 total observations.

Caus Assessment Unit / Water Name / Location Desc. Category	se ory Cause Name	Cycle First Listed	Dev.	Water Size
VAW-N16L_NEW01A02 / Claytor Lake (New River) / Claytor Lake 4C from its impounding structure upstream to the Claytor State Park Cabins.	Dissolved Oxygen		1,	196.91
VAW-N16L_NEW01B14 / Claytor Lake (New River) / Claytor Lake from the Claytor State Park Cabins upstream to the former Burlington Industries water intake.	Dissolved Oxygen			602.03
VAW-N16L_NEW02A02 / Claytor Lake (New River) / Claytor Lake 4C from the Claytor State Park Cabins upstream to the confluence of Peak Creek	Dissolved Oxygen			278.51
Claytor Lake - New River			eservoir	River
Aquatic Life		(Sq. Miles) (Acres)	(Miles)
Dissolved Oxygen - Total Impaire	ed Size by Water Type:	2	,077.45	

New River Basin

Sources:

Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

New River Basin

Cause Group Code: N16L-02-DO Claytor Lake - Peak Creek

Cause Location: Peak Creek from its confluence with the New River upstream to the end of the WQS public water supply (PWS)

designation (Dublin Quad).

City / County: Pulaski Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Dissolved Oxygen / 4C

A portion of Claytor Lake in the Peak Creek (Lower) (216.86 acres) arm is originally 2002 303(d) Listed for excursions of the Class IV Water Quality Standard (WQS) dissolved oxygen minimum criterion of 4.0 mg/l. The impairment is categorized as natural (4C) as there are no excursions of the Claytor Lake criterion for chlorophyll a (25 µg/L) or total phosphorus (20 µg/L algaecides applied) from stations 9-NEW089.34 or 9-NEW087.14 (Lacustrine zone). Virginia's Lake Nutrient Criteria (9 VAC 25-260-187) states the nutrient criteria apply only in the epilimnion for lacustrine waters during thermal stratification for control of nutrient enrichment. Guidance Memo No. 09-2005 "Monitoring and Assessment of Lakes and Reservoirs" outlines criteria for evaluating dissolved oxygen during periods of thermal stratification. Data from station 9-PKC000.00 finds the waters not supporting the Aquatic Life Use in the epilimnion from dissolved oxygen exceedances of the minimum 4.0 mg/l criterion.

9-PKC000.00 (Mouth of Peak Cr.)- The 2020 integrated Report (IR) finds 157 of 687 dissolved oxygen (DO) measurements exceed the 4.0 mg/l minimum criterion in the epilimnion. The 2018 integrated Report (IR) finds 212 of 723 dissolved oxygen (DO) measurements exceed the 4.0 mg/l minimum criterion in the epilimnion. The 2016 integrated Report (IR) finds 116 of 791 dissolved oxygen (DO) measurements exceed the 4.0 mg/l minimum criterion in the epilimnion. 2014 data report 123 of 725 DO measurements exceed the 4.0 mg/l minimum criterion in the epilimnion. 2012 DO measurements find 93 of 673 measurements in excess of the 4.0 mg/l minimum criterion. The 2010 assessment reports 69 epilimnion DO measurements exceeding the 4.0 mg/l minimum from 633 measurements. 2008 results find 131 exceed from 618 total observations.

Assessment Unit / Water Name / Location Desc.

Cause Category Cause Name

Cycle TMDL
First Dev. Water
Listed Priority Size

VAW-N17L_PKC01A10 / Claytor Lake (Peak Creek) / Peak Creek 4C Dissolved Oxygen from its confluence with the New River upstream to the end of the WQS public water supply (PWS) designation.

216.86

Claytor Lake - Peak Creek

Estuary (Sq. Miles)

Reservoir (Acres)

River (Miles)

Dissolved Oxygen - Total Impaired Size by Water Type:

216.86

Sources:

Aquatic Life

Natural Conditions - Water Quality Standards Use Attainability Analyses Needed

New River Basin

Cause Group Code: N17R-01-BAC Peak Creek and Tract Fork

Cause Location: The bacteria impairment extends from the mouth of Hogan Creek downstream to the backwaters of Claytor Lake.

And Tract Fork mainstem from its confluence with Peak Creek upstream to the mouth of Pondlick Branch.

City / County: Pulaski Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Peak Creek Bacteria Total Maximum Daily Load (TMDL) received U.S. EPA approval on 8/30/2004 [Fed. ID 7824] and SWCB approval on 12/02/2004. These waters are 1996 303(d) Listed originally for fecal coliform bacteria for 3.49 miles (4.65 mi. pre-NHD) and extended upstream in subsequent assessment cycles for a total 6.49 miles. The Recreational Use remains impaired. Tract Fork is a 2012 nested impairment within the overall Bacteria TMDL watershed. The TMDL Study can be viewed at http://www.deq.virginia.gov. Escherichia coli (E.coli) replaces fecal coliform (FC) bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters].

9-PKC011.11 (Commerce St. Bridge) Two of 11 escherichia coli (E.coli) observations exceed the WQS instantaneous criterion of 235 cfu/100 ml within the 2016 data window. Excessive values range are 325 and 350 cfu/100 ml. There are no additional E.coli data within the 2014 data window or beyond the 2008 IR. None of the three remaining samples within the 2012 data window exceed the instantaneous criterion. Data within the 2008 and 2010 data windows find two of 10 E.coli samples exceeding the 235 cfu/100 ml instantaneous criterion with exceeding values the same as in 2008. Both exceedances are 500 and 640 cfu/100 ml. E.coli results in 2006 find two of seven samples in excess of the 235 cfu/100 ml criterion; exceedances are the same as in 2008.

9-PKC009.29 (Near Radio Tower) There are no additional data beyond the 2008 IR. One exceeding value occurs within the 2012 data window at 500 cfu/100 ml of the reaming three observations. E.coli data within the 2010 data window reveal 12 exceeding values from 21 samples. The 2008 IR finds E.coli exceeds the instantaneous criterion in 12 of 23 samples. Exceeding values for both 2010 and 2008 data windows range from 240 cfu/100 ml. to 10,000. E.coli exceeds the instantaneous criterion in 11 of 18 samples in 2006 with the same range of exceedance.

9-PKC007.80 (Rt. 99 bridge) Eleven of 25 E.coli samples exceed the 235 cfu/100 ml instantaneous criterion within the 2018 data window. Excursions range from 275 to greater than 2,000 cfu/100 ml.

9-TCK000.50 (Rt. 674 Bridge)- Escherichia coli (E.coli) data within the 2012 and 2014 data windows reveal seven of 12 samples in excess of the 235 cfu/100 ml instantaneous criterion. Values in excess of the criterion range from 250 to 880 cfu/100 ml. There are no additional bacteria data within the 2016 data window.

_	Cause atego	ry Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size	
VAW-N17R_PKC01A00 / Peak Creek / This portion of Peak Creek begins just downstream of the Rt. 99/Norfolk Southern crossing extending downstream to the inundation of Peak Creek in Claytor Lake (NE46).	4A	Escherichia coli (E. coli)	2006	L	1.83	
VAW-N17R_PKC02A00 / Peak Creek / The segment begins downstream of the Washington Ave. Bridge (~0.20 miles) and extends on downstream to just below the Rt. 99 Bridge/Norfolk Southern Railway crossing of Peak Creek (NE46).	4A	Escherichia coli (E. coli)	2006	L	1.66	
VAW-N17R_PKC03A00 / Peak Creek / This portion of Peak Creek extends from the mouth of Tract Fork to downstream of the Washington Ave. Bridge (~0.20 miles) (NE46).	4A	Escherichia coli (E. coli)	2006	L	0.51	
VAW-N17R_PKC03A06 / Peak Creek / This portion of Peak Creek extends from the Magnox, Inc. outfall on downstream to the mouth of Tract Fork (NE44).		Escherichia coli (E. coli)	2006	L	0.39	
VAW-N17R_PKC04A00 / Peak Creek / The segment extends from the mouth of Hogan Creek downstream to just above the Magnox. Inc. outfall on Peak Creek (NE44).	4A	Escherichia coli (E. coli)	2006	L	2.10	

New River Basin

VAW-N17R_TCK01A00 / Tract Fork / Tract Fork mainstem from its 4A Escherichia coli (E. coli) confluence with Peak Creek upstream to the mouth of Pondlick

2012

1.24

Branch (NE45).

Peak Creek and Tract Fork Recreation

Estuary (Sq. Miles) Reservoir (Acres)

L

River (Miles)

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

7.73

Sources:

Livestock (Grazing or Feeding Operations)

Municipal (Urbanized High Density Area)

On-site Treatment Systems (Septic Systems and Similar Decentralized Systems)

Sanitary Sewer Overflows (Collection System Failures)

Unspecified Domestic

Wastes from Pets

Wildlife Other than

Waste

Waterfowl

New River Basin

Cause Group Code: N17R-01-BEN Peak Creek

Cause Location: Benthic impaired waters begin downstream of the Washington Ave. Bridge (~0.20 miles) on downstream to the

inundation of Peak Creek in Claytor Lake.

City / County: Pulaski Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

The Peak Creek General Standard - Benthic (Metals) Total Maximum Daily Load (TMDL) is U.S. EPA approved 8/30/2004 [Fed ID 7823/7822] and SWCB approval on 12/02/2004. The TMDL finds cooper (Cu) and zinc (Zn) as stressors for 3.49 miles to this 1996 (4.65 mi. pre-NHD) Listed benthic impairment. The TMDL allocations require reductions in zinc and copper from non-point sources.

9-PKC009.29 (Near Radio Tower) Bio 'IM' There are no additional data beyond the 2014 data window. Two 2011 Virginia Stream Condition Index (VSCI) surveys within the 2014 and 2016 data windows produce spring and fall scores of 31.6 and 37.4. 2011 samples show low diversity of taxa and several pollution tolerant taxa dominating the samples. Filter, collector and scraper feeding type taxa were the dominant functional feeding groups. There are no additional data within the 2010 or 2012 data windows. The 2008 IR reports four Virginia Stream Condition Index (VSCI) surveys (2002, 2003 & 2006) have an average score of 47.9. The spring 2003 sample had high diversity and numbers of mayflies compared to other samples collected in this assessment period. High flows in 2003 potentially contributed to these higher numbers. The samples with low scores show low diversity of taxa and several pollution tolerant taxa dominating the samples. Filter, collector and scraper feeding taxa were the dominant functional feeding groups. Habitat in this reach has been impacted by loss of riparian vegetation and in stream cover, and increased sedimentation.

9-PKC007.80 (Rt. 99 Bridge) Bio 'IM' Four Virginia Stream Condition Index (VSCI) surveys (2011 & 2014) within the 2016 and 2018 data windows produce an average score of 48.4. 2014 data window report impairment from two 2011 surveys. The VSCI scores are spring 33.8 and fall 58.3. Benthic community data show several pollution tolerant taxa were dominant. Mayflies typically had low abundance and other sensitive taxa such as stoneflies and caddisflies were very rare in samples. Habitat in this reach has been impacted by the loss of riparian vegetation. There are no additional data within either the 2010 or 2012 data windows. The 2008 data window reports four VSCI surveys (2002, 2003 & 2006) with an average score of 47.6. These collections reveal several pollution tolerant taxa are dominant. Habitat in this reach has been impacted by the loss of riparian vegetation.

9-PKC005.95 (Upstream of I-81 crossing)- A 2004 Probabilistic site. Two VSCI surveys, spring (62.5) and fall (58.4) result in an average score of 60.5. near the lower limit for reference conditions. Impacts from sediment deposition were noted during the spring survey. Other habitat parameters scored in the optimal to sub-optimal range. Approximately 5% of the land cover upstream of this station is urban. The TMDL study found the impairment cause to be heavy metals in sediments and storm runoff. Both samples at this station were dominated by organisms tolerant of nutrient enrichment. Since this station is within a known impaired segment and VSCI scores are near the Impaired/Non-impaired cutoff, best professional judgment designates the station as impaired.

Assessment Unit / Water Name / Location Desc.	Cause Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N17R_PKC01A00 / Peak Creek / This portion of Peak Cree begins just downstream of the Rt. 99/Norfolk Southern crossing extending downstream to the inundation of Peak Creek in Claytor Lake (NE46).	k 4A	Benthic Macroinvertebrate Bioassessments	s	1996	L	1.83
VAW-N17R_PKC02A00 / Peak Creek / The segment begins downstream of the Washington Ave. Bridge (~0.20 miles) and extends on downstream to just below the Rt. 99 Bridge/Norfolk Southern Railway crossing of Peak Creek (NE46).	4A	Benthic Macroinvertebrate Bioassessments	s	1996	L	1.66
Peak Creek Aguatic Life			Estuary Sg. Miles)		servoir .cres)	River (Miles)
Benthic Macroinvertebrates Bioassessments - Total Ir	mpaire	`	- /	ζ.	/	3.49

New River Basin

Sources:

Contaminated Sediments

Industrial/Commercial Site Stormwater Discharge (Permitted) Sediment Resuspension (Contaminated Sediment)

New River Basin

Cause Group Code: N17R-01-CU Peak Creek

Cause Location: Impairment begins downstream of the Washington Ave. Bridge (~0.20 miles) on downstream to the inundation of

Peak Creek in Claytor Lake.

City / County: Pulaski Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Copper / 4A

The Peak Creek General Standard - Benthic (Metals) Total Maximum Daily Load (TMDL) received U.S. EPA approval on 8/30/2004 [Fed ID 7823/7822] and SWCB approval on 12/02/2004.

The TMDL finds copper (Cu) and zinc (Zn) as stressors for 3.49 miles to this 1996 (4.65 mi. pre-NHD) Listed benthic impairment. The likelihood of dissolved metals reaching acute levels of toxicity in the water column during low-flow and storm events was assessed. The impact of point source discharges of Cu and Zn during low flow was analyzed and determined that the concentrations of Cu and Zn would not likely approach the acute criteria for aquatic life (i.e., 13 ug/l and 120 ug/l for Cu and Zn, respectively). It was anticipated that acidic runoff from historic industrial sites may leach significant levels of dissolved Cu and Zn to the stream during storm events. The weight of evidence at this time, including site observations and collected data, points to soils at or from the Allied Signal site as the main source of contamination.

Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N17R_PKC01A00 / Peak Creek / This portion of Peak Crebegins just downstream of the Rt. 99/Norfolk Southern crossing extending downstream to the inundation of Peak Creek in Claytor Lake (NE46).	eek 4A Copper		2006	L	1.83
VAW-N17R_PKC02A00 / Peak Creek / The segment begins downstream of the Washington Ave. Bridge (~0.20 miles) and extends on downstream to just below the Rt. 99 Bridge/Norfolk Southern Railway crossing of Peak Creek (NE46).	4A Copper		2006	L	1.66
Peak Creek		Estuary (Sg. Miles)		servoir .cres)	River (Miles)
Aquatic Life Copper - Total	Impaired Size by Water Type	, ,	(/ (3.49

Sources:

Contaminated Sediments

Industrial/Commercial Site Stormwater Discharge (Permitted)

Sediment Resuspension (Contaminated Sediment)

New River Basin

Cause Group Code: N17R-01-ZN Peak Creek

Cause Location: Impairment begins downstream of the Washington Ave. Bridge (~0.20 miles) on downstream to the inundation of

Peak Creek in Claytor Lake.

City / County: Pulaski Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Zinc / 4A

The Peak Creek General Standard - Benthic (Metals) Total Maximum Daily Load (TMDL) received U.S. EPA approval on 8/30/2004 [Fed ID 7823/7822] and SWCB approval on 12/02/2004.

The TMDL finds copper (Cu) and zinc (Zn) as stressors for 3.49 miles to this 1996 (4.65 mi. pre-NHD) Listed benthic impairment. The likelihood of dissolved metals reaching acute levels of toxicity in the water column during low-flow and storm events was assessed. The impact of point source discharges of Cu and Zn during low flow was analyzed and determined that the concentrations of Cu and Zn would not likely approach the acute criteria for aquatic life (i.e., 13 ug/l and 120 ug/l for Cu and Zn, respectively). It was anticipated that acidic runoff from historic industrial sites may leach significant levels of dissolved Cu and Zn to the stream during storm events. The weight of evidence at this time, including site observations and collected data, points to soils at or from the Allied Signal site as the main source of contamination.

Zinc - Total Im	paired Size by Water Type:				3.49
Peak Creek Aquatic Life		Estuary (Sq. Miles)		servoir cres)	River (Miles)
VAW-N17R_PKC02A00 / Peak Creek / The segment begins downstream of the Washington Ave. Bridge (~0.20 miles) and extends on downstream to just below the Rt. 99 Bridge/Norfolk Southern Railway crossing of Peak Creek (NE46).	4A Zinc		2006	L	1.66
VAW-N17R_PKC01A00 / Peak Creek / This portion of Peak Creek begins just downstream of the Rt. 99/Norfolk Southern crossing extending downstream to the inundation of Peak Creek in Claytor Lake (NE46).	4A Zinc		2006	L	1.83
	Cause ategory Cause Name		First Listed	Dev. Priority	Water Size

Sources:

Contaminated Sediments

Industrial/Commercial Site Stormwater Discharge (Permitted)

Sediment Resuspension (Contaminated Sediment)

Cycle TMDI

New River Basin

Cause Group Code: N18R-01-BAC Crab Creek

Cause Location: The upstream limit is the Crab Creek headwaters on the Ironto Quad. The downstream limit is at the Crab Creek

mouth on the New River about 1.5 mi upstream of the Rt. 114 Bridge and downstream of Radford, Virginia (Riner,

Blacksburg and Radford North Quads).

City / County: Montgomery Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Crab Creek Bacteria Total Maximum Daily Load (TMDL) is U.S. EPA approved on 8/10/2004 [Fed ID 18594 / 23405] and SWCB approved 12/02/2004 (formerly VAW-N18R-01). The waters are initially 303(d) Listed with the 2002 Assessment for fecal coliform (FC) bacteria causing non-support of the Recreational Use for 12.36 miles. The TMDL Study and allocations can be viewed at http://www.deq.virginia.gov. Escherichia coli (E.coli) replaces fecal coliform (FC) bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters].

9-CBC009.81 (Rt. 111 Bridge) There are no additional data beyond the 2010 IR where the 2010 data window finds four of 15 E.coli samples exceed the 235 cfu/100 ml instantaneous criterion. Exceedances range from 250 to greater than 2000 cfu/100 ml. Three non-exceeding E.coli observations remain within the 2012 data window. The 2008 assessment finds six of 18 E.coli samples exceed the instantaneous criterion and six of 15 exceed in 2006. The range of exceeding values is from 400 to greater than 2000 cfu/100 ml in 2008 and 2006.

9-CBC006.35 (Rt. 661 Bridge) Both the 2010 and 2012 data windows find four of 12 E.coli samples exceeding the instantaneous criterion. The range of exceedance is from 380 to 950 cfu/100 ml. E.coli data within the 2008 data window are three of six exceeding values. The 2006 assessment reports E.coli exceeds the WQS instantaneous criterion of 235 cfu/100 ml in eight of 16 observations. Exceeding values range from 250 to >800 cfu/100 ml. This station is located upstream of the former Christiansburg outfall.

9-CBC004.38 (Rt. 660 Bridge) There are no additional data beyond the 2010 data window. Five of 15 remaining escherichia coli (E.coli) observations in 2012 exceed the 235 cfu/100 ml instantaneous criterion ranging from 250 to 1200 cfu/100 ml. Data within the 2010 data window find exceedances ranging from 250 to 1200 cfu/100 ml in 14 of 35 observations. E.coli exceeds the 235 cfu/100 ml WQS instantaneous criterion in 16 of 33 observations within the 2008 data window. Exceeding values range from 280 to greater than 800 cfu/100 ml. 2006 E.coli results find 22 of 40 observations in excess of the instantaneous criterion and the same range of exceedance.

9-CBC001.00 (Route 663 Bridge near Walton) The 2020 data window finds five of 12 excursions. There are no additional data beyond the 2014 IR where six of 24 E.coli observations exceed the instantaneous criterion ranging from 250 to greater than 2000 cfu/100 ml. There were no additional data within the 2010 and 2012 data windows. Two of 15 remaining E.coli observations in 2012 exceed the instantaneous criterion at 250 and 1300 cfu/100 ml. 2010 values exceeding the instantaneous criterion range from 250 to 1300 cfu/100 ml in 10 of 35 samples. Nine of 27 E.coli samples exceed the instantaneous criterion ranging from 260 to greater than 800 cfu/100 ml in 2008. The 2006 Integrated Report (IR) finds nine of 23 E.coli samples exceed the instantaneous criterion. The range of exceeding values is the same as in 2008.

_	Cause Catego	e ry Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size	
VAW-N18R_CBC01A00 / Crab Creek / This section of the mainstem Crab Creek extends from its mouth on the New River on upstream of the Walton community (NE58).	4A	Escherichia coli (E. coli)	2004	L	2.15	
VAW-N18R_CBC02A00 / Crab Creek / These mainstem waters of Crab Creek extend from upstream of the Walton community to upstream of the Vicker community. The end of the WQS public wate supply (PWS) designation (NE58).		Escherichia coli (E. coli)	2004	L	1.18	
VAW-N18R_CBC03A00 / Crab Creek / These waters are the Crab Creek mainstem from upstream of the Vicker community on upstream to the former Christiansburg STP outfall on Crab Creek (NE58).	4A	Escherichia coli (E. coli)	2004	L	1.10	

New River Basin

VAW-N18R_CBC04A00 / Crab Creek / These mainstem waters extend from the former Christiansburg STP outfall upstream to Crab Creek's headwaters (NE58).

4A Escherichia coli (E. coli)

2004

7.93

Crab Creek

Recreation

Estuary (Sq. Miles)

Reservoir (Acres)

L

River (Miles)

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

12.36

Sources:

Discharges from Municipal Separate Storm Sewer Systems (MS4) Livestock (Grazing or Feeding Operations)

Municipal (Urbanized High Density Area)

Unspecified Domestic Waste

Wastes from Pets

Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N18R-01-BEN Crab Creek

Cause Location: The upstream limit is the Crab Creek headwaters on the Ironto Quad. The downstream limit is at the Crab Creek

mouth on the New River about 1.5 mi upstream of the Rt. 114 Bridge and downstream of Radford, Virginia (Riner,

Blacksburg and Radford North Quads).

City / County: Montgomery Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

The 1996 303(d) Listing of the Crab Creek General Standard (Benthic) Total Maximum Daily Load (TMDL) Study is U.S. EPA approved 8/10/2004 [Sediment- Fed ID 18595 / 23406]. The SWCB approved the TMDL on 12/02/2004 (formerly VAW-N18R-01). The TMDL identifies sediment to be the primary stressor, with organic matter and nutrient enrichment as additional stressors. The waters remain impaired for the aquatic life use for 12.36 miles.

Natural seasonal effects are noted at the sites below. Pollution tolerant families are dominant in both seasons, the midge family Chironomidae in spring and the caddisfly family Hydropsychidae in fall. Beginning in spring 2002, Toms Creek was determined to be a more suitable ecoregion reference site because of similarity in size and watershed characteristics than the previous reference site (Sinking Creek, 9-SNK012.06). Agricultural and urban NPS runoff impact Crab Creek. Habitat impacts to this reach result in fine sediment deposition causing stream substrates to become embedded from bank erosion, altered hydrology, and degraded riparian buffers due to residences, roads, and railroad tracks. An apparent nutrient rich environment all contribute to the benthic impairment.

9-CBC007.55- Bio 'IN' with one sample collected in 2013 to evaluate effect of tributary stream restoration. VSCI score of 37.2

9-CBC006.35- Bio 'IM'; No additional data beyond the 2010 data window. Two 2008 Virginia Stream Condition Index (VSCI) surveys with an average score of 39.36 are within the 2012 and 2014 data windows. Three VSCI surveys (2003 & 2008) result in an average VSCI score of 43.33 are within the 2010 data window. Moderately pollution tolerant to pollution tolerant organisms (oligochaetes, chironomidae, hydropsychidae, and elmidae) are dominant in both seasons. Habitat impacts to this reach result in fine sediment deposition that causes stream substrates to become embedded, altered hydrology, and degraded riparian buffers due to roads. The 2008 IR reports five RBP II surveys scoring- 2000 spring 47.83, fall- 34.78; 2002 spring-52.17, fall- 59.09 and 2003 spring- 65.22. Seasonal 5 year Spring score 55.07 and Fall score 46.94.

9-CBC004.38- Bio 'IM'; No additional data beyond the 2010 data window. Two 2008 VSCI surveys lie within the 2012 and 2014 data window with an average score of 53.0. Three VSCI surveys (2003 & 2008) with an average score of 52.28 are produced within the 2010 data window. There is some difference in the biological condition scores between seasons. Fall samples showed an increase of %mayflies over the spring samples. Agricultural and urban NPS runoff impact Crab Creek. Habitat impacts to this reach result in fine sediment deposition that causes stream substrates to become embedded, bank erosion from altered hydrology, and degraded riparian buffers due to pastures and railroad tracks. Five RBP II surveys scoring- 2000 spring-39.13, fall- 34.78; 2002 spring- 65.22, fall- 59.09 and 2003 spring- 69.57. Seasonal 5 year Spring score 57.97; Fall score 46.94 are reported in the 2008 IR.

9-CBC001.00- No additional data beyond the 2014 data window. Bio 'IM' The 2012 and 2014 data windows find an average score of 55.0 from two surveys (2008). The 2010 IR finds impairment remains from three VSCI surveys (2003 & 2008) with an average score of 60.0. The moderately pollution tolerant midge family Chironomidae is dominant in both seasons. Impacts to the benthos and stream habitat are the same as noted at 9-CBC004.38. The 2008 IR reports three VSCI surveys (2002-2003) with an average score of 58.43. Pollution tolerant families are dominant in spring and fall, the midge family Chironomidae in spring and the caddisfly family Hydropsychidae in fall. Impacts to the benthic community and stream habitat are the same as noted at 9-CBC004.38.

Cycle TMDL

Assessment Unit / Water Name / Location Desc.	Cause Catego	e rry Cause Name	First Listed	Dev. Priority	Water Size
VAW-N18R_CBC01A00 / Crab Creek / This section of the mainstem Crab Creek extends from its mouth on the New River on upstream of the Walton community (NE58).	4A	Benthic Macroinvertebrates Bioassessments	1996	L	2.15
VAW-N18R_CBC02A00 / Crab Creek / These mainstem waters Crab Creek extend from upstream of the Walton community to	of 4A	Benthic Macroinvertebrates Bioassessments	1996	L	1.18

New River Basin

upstream of the Vicker community. The end of the WQS public water supply (PWS) designation (NE58).

VAW-N18R CBC03A00 / Crab Creek / These waters are the Crab Creek mainstem from upstream of the Vicker community on upstream to the former Christiansburg STP outfall on Crab Creek (NE58).

VAW-N18R_CBC04A00 / Crab Creek / These mainstem waters extend from the former Christiansburg STP outfall upstream to Crab Creek's headwaters (NE58).

Benthic Macroinvertebrates Bioassessments

Estuary

(Sq. Miles)

1996 1.10

L

Reservoir

Benthic Macroinvertebrates Bioassessments

1996

7.93

River

Crab Creek

Aquatic Life

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type:

(Acres) (Miles) 12.36

Sources:

Discharges from Municipal Separate Storm Sewer Systems (MS4)

Sediment Resuspension

(Clean Sediment)

Loss of Riparian Habitat

Sediment Resuspension (Contaminated Sediment) Municipal (Urbanized High

Density Area)

Streambank Modifications/Destabilization Post-development Erosion and Sedimentation

New River Basin

Cause Group Code: N18R-02-BAC **Connellys Run**

Cause Location: Bacteria impairment begins near the headwaters of Connellys Run at an unnamed tributary (37°07'04" / 80°32'16")

downstream to its mouth on the New River.

City / County: Radford City

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

Fecal coliform (FC) bacteria excursions of the former WQS 400 cfu/100 ml instantaneous criterion cause non-support of the Recreational Use for 2.85 miles. The impairment for the 2004 303(d) Listed water remains. Escherichia coli (E.coli) replaces fecal coliform (FC) bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170, Bacteria: other waters].

9-CNL000.01 (Bissett Park Bridge, Radford) One of six E.coli excursions reported during the 2020 data window. There are no additional data since the 2014 IR where escherichia coli (E.coli) data exceed the 235 cfu/100 ml instantaneous criterion in five of 24 samples. Excessive values range from 790 to greater than 2000 cfu/100 ml. 2012 and 2010 E.coli data exceed the 235 cfu/100 ml instantaneous criterion in four of 12 samples. Excessive values range from 260 to 1260 cfu/100 ml. The 2006 assessment finds FC exceedances of the former WQS instantaneous criterion of 400 cfu/100 ml in three of 11 observations. The range of excursions is from 500 to 1900 cfu/100 ml. The initial 2004 303(d) Listing is based on FC exceedances of the former WQS instantaneous criterion of 400 cfu/100 ml in three of nine observations with the range of exceedance the same as 2006.

90CNL000.06 - This station is not the 303(d) listing station but reports three of 6 E.coli excursions during the 2020 data window.

Assessment Unit / Water	Name / Location Desc.	Cause Category Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
unnamed tributary @37°07'23"	nnellys Run / Connellys Run from /80°33'21"; 1.57 miles upstream nstream to its confluence on the N	of)	2010	Н	1.60
near Rt. 611 @ 37°07'04" / 80°3	eam to the confluence of an unnar 1"; 1.57 miles upstream of the)	2010	Н	1.25
Connellys Run			Estuary		servoir	River
Recreation			(Sq. Miles)	(A	cres)	(Miles)
	Escherichia coli (E. coli) - Tota	al Impaired Size by Water Type:				2.85
Sources:						
Livestock (Grazing or Feeding Operations)	Municipal (Urbanized High Density Area)	Unspecified Domestic Waste	Wastes from Pets			
Wildlife Other than Waterfowl						

New River Basin

Cause Group Code: N19R-01-BAC Little River (Upper)

Cause Location: The bacteria impaired waters begin in the headwaters of Little River and extend downstream to the mouth of the West Fork of Little River (Check, Endicott and Flovd Quads).

City / County: Floyd Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013. The original 2004 fecal coliform (FC) bacteria 303(d) Listing is extended downstream and upstream based on escherichia coli (E.coli) bacteria collections within the 2006 data window. The waters are impaired for 34.67 miles for failure to support the Recreational Use. Escherichia coli (E.coli) replaces fecal coliform (FC) bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters].

9-LRV069.88 (Rt. 641 Bridge) There are no additional data beyond the 2008 Integrated Report (IR) where four of 12 E.coli observations exceed the 235 cfu/100 ml instantaneous criterion within the 2008 and 2010 data windows. Values in excess of the criterion range from 500 to 1500 cfu/100 ml. The 2006 IR reports three of nine E.coli observations exceed the instantaneous criterion. Values in excess of the criterion range from 350 to 1500 cfu/100 ml.

9-LRV065.57 (Rt. 639 Bridge) Seven of 12 E.coli samples exceed the 235 cfu/100 ml instantaneous criterion within the 2018 data window. Excursions range from 266 to 805 cfu/100 ml. The 2012 data window finds E.coli exceeds the instantaneous criterion in seven of 15 samples. Exceeding values range from 350 to 1800 cfu/100 ml. Both the 2008 and 2010 data windows find escherichia coli (E.coli) exceeds the instantaneous criterion in four of 11 samples. Exceeding values range from 430 to 800 cfu/100 ml.

9-LRV056.74 (Rt. 221 Bridge) There are no additional data beyond the 2008 IR where four of 12 E.coli observations exceed the 235 cfu/100 ml instantaneous criterion. Maximum values exceeding the criterion range from 400 cfu/100 ml to greater than 2000. The 2006 assessment finds three of nine E.coli observations exceed the instantaneous criterion with the same range of exceedance as 2008. The original 2004 303(d) Listing is based on exceedance of the former fecal coliform bacteria 400 cfu/100 ml instantaneous criterion where two observations exceed from 11 samples. 2004 IR FC values exceeding the standard are 500 and 1400 cfu/100 ml.

9-LRV044.49 (Rt. 615 Bridge)2020 data window: one of 12 excursions. There are no additional data beyond the 2014 data window. Escherichia coli (E.coli) exceedances are found in four of 12 observations. Exceeding values range from 1650 to greater than 2000 cfu/100 ml within the 2014 data window. There are no additional data within the 2010 and 2012 data windows. The 2008 IR reports E.coli exceedances are found in three of 11 observations. Exceeding values range from 380 to greater than 2000 cfu/100 ml. Two of eight E.coli exceedances are found in 2006 at 380 and 450 cfu/100 ml.

Assessment Unit / Water Name / Location Desc.	Cause Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N19R_LRV01A00 / Little River / Little River mainstem waters from the West Fork Little River confluence upstream to the mouth of Oldfield Creek (NE49).		Escherichia coli (E. coli)		2006	L	8.73
VAW-N19R_LRV02A00 / Little River / Little River mainstem waters from the mouth of Oldfield Creek upstream to the mouth of Beaverdam Creek (NE49).	s 4A	Escherichia coli (E. coli)		2006	L	7.59
VAW-N19R_LRV03A00 / Little River / Little River mainstem waters from the mouth of Beaverdam Creek upstream to near its headwaters (NE48).	s 4A	Escherichia coli (E. coli)		2006	L	18.35
Little River (Upper)			Estuary	Res	servoir	River
Recreation			(Sq. Miles)	(A	cres)	(Miles)
Escherichia coli (E. coli) - Total Im	npaire	d Size by Water Type:				34.67

New River Basin

Sources:

Livestock (Grazing or Feeding Operations)

Wildlife Other than Waterfowl

On-site Treatment Systems (Septic Systems and Similar Decentralized Systems) Unspecified Domestic Waste

Wet Weather Discharges (Non-Point Source)

New River Basin

Cause Group Code: N19R-01-TEMP Little River

Cause Location: Little River mainstem waters from the mouth of the West Fork Little River upstream to the mouth of Payne Creek.

City / County: Floyd Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 4A

The Little River Temperature (Fed ID: 41518) TMDL Study U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013. The 303(d) Listed natural trout water temperature impairment is extended both upstream and downstream in 2008 from the original impairment defined by station 9-LRV056.74 in 2002. The upstream extension is based on station 9-LRV065.57. And the downstream extension on station 9-LRV044.49. Total non-support of the Aquatic Life Use is 34.67 miles.

9-LRV065.57- (Rt. 639 Bridge) The Class VI 20°C criterion is exceeded within the 2018 data window at 23.4°C (6/22/15) and 22.3°C (9/2/15). The 2012 data window reports temperature exceedances in two of 19 measurements. Excursions are 20.4 °C on 6/29/2005 and 23.7°C on 8/5/2010. Temperature exceedances are found in two of 12 measurements in 2008 and 2010. Each are in excess of the WQS Class VI natural trout water criterion of 20°C. Excursions are both at 20.4 °C on 8/02/2004 and 6/29/2005.

9-LRV056.74- (Rt. 221 Bridge) Temperature data within the 2014 data window are insufficient to de-list these waters (0/4 samples). The temperature impairment remains. The 2008 IR reports temperature exceedances of the natural trout water criterion occur in two of 12 measurements. The excursions occur on 8/02/2004 at 21.4 °C and 6/29/2005 at 21.3°C within the 2008 data window. The 2006 Integrated Report (IR) records two of 12 temperature measurements exceeding the criterion with excursions in May of 2000 (at 21.2°C) and August of 2004 (at 21.4 °C). Two of 11 measurements exceed in 2004. The exceedances occur in July 1998 (at 25.7°C) and May of 2000 (at 21.2°C). The 2002 assessment found temperature exceeds in three of 16 measurements occurring in July 1997 and 1998 (2) and one in May of 2000.

9-LRV044.49- (Rt. 615 Bridge) There are no additional data beyond the 2014 data window. Three temperature measurements exceed the 20°C natural trout criterion at 26.2°C (7/21/2011), 20.5°C (9/13/2011) and 23.9°C (8/29/2012) from 12 measurements within the 2014 data window. There were no additional data within the 2010 and 2012 data windows. The 2008 IR reports two temperature measurements exceed the Class VI 20 °C natural trout criterion at 23.3 °C (8/02/2004) and 22.8°C (6/29/2005) from 12 measurements.

Assessment Unit / Water Name / Location Desc.	Caus Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N19R_LRV01A00 / Little River / Little River mainstem wate from the West Fork Little River confluence upstream to the mouth o Oldfield Creek (NE49).		Temperature		2008	L	8.73
VAW-N19R_LRV02A00 / Little River / Little River mainstem wate from the mouth of Oldfield Creek upstream to the mouth of Beaverdam Creek (NE49).	rs 4A	Temperature		2002	L	7.59
VAW-N19R_LRV03A00 / Little River / Little River mainstem wate from the mouth of Beaverdam Creek upstream to near its headwaters (NE48).	rs 4A	Temperature		2008	L	18.35
Little River			Estuary	Res	servoir	River
Aquatic Life			(Sq. Miles)) (A	cres)	(Miles)
Temperature - Total I	mpaire	d Size by Water Type:				34.67

Sources:

Loss of Riparian Habitat Natural Sources

New River Basin

Cause Group Code: N19R-02-BAC Meadow Run

Cause Location: Meadow Run (MDR) from its headwaters downstream to its confluence with Little River.

City / County: Floyd Co.
Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Recreational Use remains impaired for 4.00 miles for the original 2006 303(d) Listing. The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013.

9-MDR000.34 (Rt. 641 Bridge) There are no additional data beyond the 2012 Integrated Report (IR) where nine of 15 Escherichia coli (E.coli) samples exceed the 235 cfu/100 ml instantaneous criterion. Exceeding values range from 400 to 1200 cfu/100 ml. The 2008 and 2010 IRs report four of 12 E.coli samples exceed the 235 cfu/100 ml instantaneous criterion. Exceeding values range from 630 to greater than 2000 cfu/100 ml. The 2006 range of exceedance is the same from three of nine E.coli observations.

Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N19R_MDR01A04 / Meadow Run / Meadow Run from its headwaters downstream to its confluence with Little River (NE48).	• • • • • • • • • • • • • • • • • • • •		2006	L	4.00
Meadow Run Recreation		Estuary (Sq. Miles)		servoir cres)	River (Miles)
Escherichia coli (E. coli) - Tota	I Impaired Size by Water Type:				4.00

Sources:

Livestock (Grazing or Feeding Operations)

On-site Treatment Systems

Unspecified Domestic Wet Weather Discharges
(Septic Systems and Similar Waste
Decentralized Systems)

Wet Weather Discharges
(Non-Point Source)

Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N19R-02-BEN Meadow Run

Cause Location: Meadow Run (MDR) from its headwaters downstream to its confluence with Little River.

City / County: Floyd Co.
Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

The Little River Benthic (Sediment Fed ID: 41517) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013. The original 2008 assessment finds the Aquatic Life Use impaired for 4.00 miles from the results of Virginia Stream Condition Index (VSCI) surveys.

9-MDR003.60 (Off Rt. 610) Bio 'IM' There are no additional data beyond the 2008 IR where two 2001 VSCI surveys with an average score of 45.8 are reported. The benthic community was considerably better in the fall (score 60.6) although taxa richness and percentage of stoneflies-caddisflies (Hydropsychidae) were still low. The station is located downstream and adjacent to residences with mowed lawns, a driveway and a horse pasture that impact bank vegetation and the riparian zone in this reach. The stream substrate is impacted by sediment deposition.

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type:					
Meadow Run Aquatic Life			eservoir Acres)	River (Miles)	
VAW-N19R_MDR01A04 / Meadow Run / Meadow Run from its headwaters downstream to its confluence with Little River (NE48).	4A Benthic Macroinvertebrates Bioassessments	2008	L	4.00	
Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	First Listed		Water Size	

Sources:

Loss of Riparian Habitat Sediment Resuspension Streambank (Clean Sediment) Modifications/Destabilization

New River Basin

Cause Group Code: N19R-03-BAC Pine Creek

Cause Location: Pine Creek mainstem from its mouth on Little River upstream to the impounding structure of a pond.

City / County: Floyd Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013. The waters remain impaired for non-support of the Recreational Use. Bacteria exceedances cause the 2006 303(d) Listing for 3.91 miles.

9-PNC000.69 (Rt. 682 Bridge) There are no additional data beyond the 2008 IR where escherichia coli (E.coli) exceed the 235 cfu/100 ml instantaneous criterion in three of 11 samples in 2008. Excursions range from 380 to 1000 cfu/100 ml. 2006 E.coli exceedances are three of eight with the same range of exceedance found in 2008.

Cause Assessment Unit / Water Name / Location Desc. Cause Category Cause Name	Cycle TMDL First Dev. Wa Listed Priority Si	ater ize
VAW-N19R_PNC01A06 / Pine Creek / Pine Creek mainstem from 4A Escherichia coli (E. coli) its mouth on Little River upstream to just above the intersection of Sandy Flats Road (Rt. 690) (NE49).	2006 L 3.	.91

Pine Creek		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			3.91

Sources:

Livestock (Grazing or Feeding Operations)

On-site Treatment Systems

Unspecified Domestic Wet Weather Discharges
(Septic Systems and Similar Waste
Decentralized Systems)

Wet Weather Discharges
(Non-Point Source)

Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N19R-03-TEMP **Pine Creek**

Cause Location: Pine Creek mainstem from its mouth on Little River upstream to the impounding structure of a pond.

City / County: Floyd Co. Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 4A

The Little River Temperature (Fed ID: 41518) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013. The Aquatic Life Use is not supported due to temperature exceedances of the WQS Class VI natural trout water criterion. The impairment extends 3.91 miles.

9-PNC000.69- There are no additional data beyond the 2008 IR. Two of 12 temperature measurements exceed the natural trout water criterion of 20°C. Each excursion is 20.5 °C on 8/02/2004 and 21.3°C on 6/29/2005.

		Cycle	TMDL	
	Cause	First	Dev.	Water
Assessment Unit / Water Name / Location Desc.	Category Cause Name	Listed	Priority	Size
VAW-N19R_PNC01A06 / Pine Creek / Pine Creek mainste	m from 4A Temperature	2008	L	3.91
its mouth on Little River upstream to just above the intersection	on of			
0				

Sandy Flats Road (Rt. 690) (NE49).

Pine Creek Estuary River Reservoir (Sq. Miles) (Acres) (Miles) **Aquatic Life** Temperature - Total Impaired Size by Water Type: 3.91

Sources:

Loss of Riparian Habitat **Natural Sources**

New River Basin

Cause Group Code: N20R-01-BAC Dodd Creek and West Fork Dodd Creek

Cause Location: Dodd Creek: The upper limit extends from the junction of Routes 710 and 714 downstream to the Dodd Creek mouth on the West Fork Little River (Woolwine and Floyd Quads).

> West Fork Dodd Creek and unnamed tributary XDC: Mainstem extends from its confluence with Dodd Creek upstream to the mouth of an unnamed tributary (XDC). The mouth of the unnamed tributary is located at 36°52'33" /

West Fork Little River: West Fork Little River waters from its mouth on Little River upstream to the mouth of Dodd Creek on the West Fork Little River.

City / County: Floyd Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

Fecal Coliform / 4A

The Dodd Creek Bacteria Total Maximum Daily Load (TMDL) is U.S. EPA approved 12/11/2002 [Fed ID 9456/23407] and SWCB approved 6/17/2004 (formerly VAW-N20R-01). The Bacteria Implementation Plan (IP) received SWCB approval 6/27/2007. The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved 3/14/2012 and SWCB approved 3/25/2013. The waters were originally 1998 303(d) listed based on the former fecal coliform (FC) WQS instantaneous criterion of 1000 cfu/100 ml and 200 geometric mean (8.90 mi). Escherichia coli (E.coli) replaces FC bacteria as the indicator as per WQS [9 VAC 25-260-170. Bacteria; other waters]. Additional bacteria sampling above and below the 1998 303(d) Dodd Creek Impaired waters have extended the original size. Tributary additions include West Fork Dodd Creek (7.04 mi) and an unnamed tributary (XDC) in 2002 to the West Fork (0.53 mi).

Dodd Creek:

9DDD-1-NCNR- Citizen Ly. 2 data for E.coli find a 'High' probability of adverse conditions from 5/8 samples ranging from 350-1400 cfu/100 ml. There are no additional data beyond the 2012 IR.

9-DDD004.64 (Rt. 720 Br above Floyd STP) No additional data beyond 2008 IR:2/9 at 280 & 1200 cfu/100 ml. 2004: 3/11 FC samples exceed the former WQS 400 cfu/100 ml instantaneous criterion.

9-DDD002.62 (Rt. 696 Br below Floyd STP) No additional data beyond 2014 IR: 21/36 E.coli samples exceed ranging from 250 to >2,000 cfu/100 ml. 2012: 21/33, 2010: 15/21 and 2008: 6/9 with same exceedance range as 2014.

9-DDD001.00 (Rt. 8 Br below Floyd STP) No additional data beyond 2014 IR: 16/36 from 250 to >2,000 cfu/100 ml. 2012: 13/33 from 250 to >2,000 cfu/100 ml. 2010: 9/21 from 350 to >2,000 cfu/100 ml. 2008: 2/9 at 350 and 1,900 cfu/100 ml.

9-DDD008.20- No additional data beyond 2004 IR: 3/3 FC exceedances of the former 400 cfu/100 ml WQS instantaneous criterion (max. 1700); 1 FC geomean calc results in the exceedance of the former 200 cfu/100 ml standard. No E.coli samples collected.

West Fork Dodd Creek:

9-DDW004.02 (Rt. 714 Br) No additional data beyond 2004 IR: 4/4 FC exceedances of the former WQS 400 cfu/100 ml instantaneous criterion (max. 9200).

9-DDW000.02- (Rt. 8 Br) No additional data beyond 2014 IR: 20/36 E.coli samples exceed ranging from 250-1800 cfu/100 ml. 2012: 12/24 from 250-1800 cfu/100 ml. 2010: 7/12 from 250-1600 cfu/100 ml.

Unnamed Tributary XDC: (The unnamed tributary portion extends from just upstream of the Rt. 8 crossing downstream to its confluence with West Fork Dodd Creek - Floyd Quad.)

9-XDC000.48 (Rt. 807 Br) No additional data beyond 2004 IR: 4/4 FC exceedances of the former WQS 400 cfu/100 ml instantaneous criterion (max. 6400).

		Cycle	TMDL	
	Cause	First	Dev.	Water
Assessment Unit / Water Name / Location Desc.	Category Cause Name	Listed	Priority	Size
VAW-N20R_DDD01A00 / Dodd Creek / Dodd Creek mainstem	4A Escherichia coli (E. coli)	2008	L	3.84

Final 2020 Appendix 5 - 3453

waters from its mouth on the West Fork of Little River upstream to the Floyd/Floyd County PSA outfall on Dodd Creek (NE51).

New River Basin							
waters from the Floyd/Floyd Cou	odd Creek mouth on Dodd Creek,	4A	Escherichia coli (E. coli))	2008	L	2.60
VAW-N20R_DDW01A02 / Wes Dodd Creek mainstem from its c upstream to the mouth of an unn the unnamed tributary is located	onfluence with Dodd Creek named tributary (XDC). The mouth	4A of	Escherichia coli (E. coli))	2010	L	1.31
Dodd Creek and West Fork Do	odd Creek			Estuary	Res	servoir	River
Recreation				(Sq. Miles)	(A	cres)	(Miles)
	Escherichia coli (E. coli) - Total	Impaired	d Size by Water Type:				7.75
Assessment Unit / Water N	ame / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
from the West Fork of Dodd Cree	upstream near the junction of Rout	4A tes	Fecal Coliform		1998	L	2.46
VAW-N20R_DDW02A02 / Wes Dodd Creek mainstem from the (XDC) upstream to its headwater tributary is located @36°52'33" /	confluence of an unnamed tributar rs. The mouth of the unnamed	4A y	Fecal Coliform		1998	L	5.73
unnamed tributary (XDC) to the	waters. The mouth of the unname		Fecal Coliform		2002	L	0.53
Dodd Creek and West Fork Do	odd Creek			Estuary	Res	servoir	River
Recreation				(Sq. Miles)	(A	cres)	(Miles)
	Fecal Coliform - Total	Impaired	d Size by Water Type:				8.72
Sources:							
Livestock (Grazing or	On-site Treatment Systems	Unspe	cified Domestic	Wet W	eather I	Discharge	es
Feeding Operations)	(Septic Systems and Similar Decentralized Systems)	Waste			oint So	•	
Wildlife Other than							

Final 2020 Appendix 5 - 3454

Waterfowl

New River Basin

Cause Group Code: N20R-01-TEMP West Fork Dodd Creek

Cause Location: West Fork Dodd Creek mainstem from its confluence with Dodd Creek upstream to the mouth of an unnamed

tributary (XDC). The mouth of the unnamed tributary is located @36°52'33" / 80°19'43".

City / County: Floyd Co.
Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 4A

The Little River Temperature (Fed ID: 41518) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board

(SWCB) approved 3/25/2013.

9-DDW000.02 (Rt. 807 Bridge) There are no additional data beyond the 2014 data window. 2014 data reveal five of 36 temperature measurements in excess of the WQS Class VI 20°C criterion. Temperature exceedances in addition to those within the 2012 IR are 24.6°C on 7/21/2011 and 22.2°C on 8/29/2012. 2012 Class VI temperature exceedances are found in three of 24 measurements occurring on 7/18/2007 at 20.9°C: 9/11/2007 at 22.3°C and 24.3°C on 8/5/2010. Temperature exceedances within the 2010 data window are found in two of 12 measurements that occur on 7/18/2007 at 20.9°C and 9/11/2007 at 22.3°C. 2002 IR reports temperature exceeds the 20° natural trout criterion in two of two measurements. Exceeding values are 23.3°C on 7/28/99 and 20.1°C on 6/28/00. The 2002 Temperature 303(d) Listing remains.

Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	First Listed	Dev. Priority	Water Size
VAW-N20R_DDW01A02 / West Fork Dodd Creek / West Fork Dodd Creek mainstem from its confluence with Dodd Creek upstream to the mouth of an unnamed tributary (XDC). The mouth	4A Temperature	2002	L	1.31
the unnamed tributary is located @36°52'33" / 80°19'43" (NE51).				

West Fork Dodd Creek

Aquatic Life

Estuary Reservoir River (Sq. Miles) (Acres) (Miles)

Temperature - Total Impaired Size by Water Type:

1.31

Cyala

TMDI

Sources:

Natural Sources

New River Basin

Cause Group Code: N20R-02-TEMP **Dodd Creek**

Cause Location: Dodd Creek from it's confluence with the West Fork Little River upstream to the mouth of the West Fork of Dodd

Creek

City / County: Floyd Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 4A

There are no additional Dodd Creek data beyond the 2014 assessment. The 2012 assessment finds the Aquatic Life Use is impaired for 8.90 miles due to temperature exceedances of these Class V (21°C) stockable trout waters criterion. The impairment is extended upstream 2.19 miles with citizen data from station 9DDD-1-NCNR in the 2010 assessment. The Little River Temperature (Fed ID: 41518) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013.

Dodd Creek (Lower): Length 3.84 miles.

9-DDD002.62- (Route 696 Bridge below Floyd STP) There are no additional data beyond the 2014 data window. The 2014 Integrated Report (IR) finds the 21°C stockable trout water criterion exceeds in four of 36 measurements at 21.7°C on 9/11/2007; 23.9°C on 8/5/2010; 23.9°C on 7/21/2011 and 21.6°C on 8/29/2012. The 21°C Class V criterion exceeds in four of 33 measurements at 22.2°C on 8/10/2005; 21.6°C on 8/14/2006; 21.7°C on 9/11/2007 and 23.9°C on 8/5/2010 within the 2012 data window. The 2010 IR finds three of 21 measurements at 22.2°C on 8/10/2005; 21.6°C on 8/14/2006; and 21.7°C on 9/11/2007. 2008 results report two of nine measurements at 22.2°C on 8/10/2005 and 21.6°C on 8/14/2006.

9-DDD001.00- (Route 8 Bridge below Floyd STP)There are no additional data beyond the 2014 data window. The 2014 Integrated Report (IR) finds the stockable trout water criterion exceeds in four of 36 measurements at 21.1°C on 9/11/2007: 23.7°C on 8/5/2010; 24.4°C on 7/21/2011 and 21.9°C on 8/29/2012. The 2012 IR reports the Class V criterion exceeds in four of 33 measurements at 22.0 on 8/10/2005; 22.1°C on 8/14/2006; 21.1°C on 9/11/2007 and 23.7°C on 8/5/2010. The 2010 assessment finds the stockable trout water criterion exceeds in three of 21 measurements at 22.0 on 8/10/2005; 22.1°C on 8/14/2006; and 21.1°C on 9/11/2007. The 2008 IR found two of nine temperature measurements exceed the Class V criterion at 22.0 on 8/10/2005 and 22.1°C on 8/14/2006.

Dodd Creek (Upper) Length 5.06 miles.

9DDD-1-NCNR (Rt. 710 Bridge) There are no additional data beyond the 2012 IR where Citizen Level 3 data finds three of 14 temperature measurements exceed the Class V criterion of 21°C. Excessive values are 25°C on 6/8/2008; 22.5°C on 8/10/2008; and 22.5°C on 9/14/2008. The 2010 data window reveals three of eight temperature measurements exceeding the criterion on the same dates in 2010. These data extended the temperature impairment upstream 2.19 miles in 2010.

Single measurement exceedances of the Class V criterion occur upstream in 2008 and 2010. There are no additional data reported for Station 9-DDD004.64 (Rt. 720 Bridge above Floyd STP) where one temperature exceedance from nine measurements is found at 22.4°C on 8/10/2005 within the 2008, 2010 and 2012 data windows.

Historically stations 9-DDD006.27 (Rt. 8 Bridge), 9-DDD004.75 (Rt. 720 Bridge) and 9-DDD004.64 (Route 720 Bridge above Floyd STP) have recorded temperature excursions upstream albeit in drought conditions. 9-DDD006.27 21.6°C on 7/28/99 -One of two temperature measurements exceed the 21°C criterion. 9-DDD004.75 records one excursion at 21.9°C on 7/28/99. The extension of the impairment to the mouth of the West Fork of Dodd Creek is in recognition of these data and temperature exceedances on the West Fork of Dodd Creek.

Cycle TMDL

Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	First Listed	Dev. Priority	Water Size	
VAW-N20R_DDD01A00 / Dodd Creek / Dodd Creek main waters from its mouth on the West Fork of Little River upstrethe Floyd/Floyd County PSA outfall on Dodd Creek (NE51).	·	2008	L	3.84	
VAW-N20R_DDD02A00 / Dodd Creek / Dodd Creek mair waters from the Floyd/Floyd County PSA outfall on Dodd Crupstream to the West Fork of Dodd Creek mouth on Dodd C just upstream of the Rt. 8 Bridge (NE51).	eek	2008	L	2.60	
VAW-N20R_DDD03A02 / Dodd Creek / Dodd Creek main	nstem 4A Temperature	2010	L	2.46	
Final 2020	Appendix 5 - 3456				

New River Basin

from the West Fork of Dodd Creek mouth on Dodd Creek, just upstream of the Rt. 8 Bridge on upstream near the junction of Routes 710 and 714 near the Blue Ridge Parkway (NE51).

Dodd Creek

Aquatic Life

Temperature - Total Impaired Size by Water Type:

Estuary (Sq. Miles)

Reservoir (Miles)

River (Miles)

8.90

Sources:

Natural Sources

New River Basin

Cause Group Code: N20R-03-TEMP West Fork Little River

Cause Location: West Fork Little River waters from its mouth on Little River upstream to the mouth of Dodd Creek on the West Fork

Little River (NE51).

City / County: Floyd Co. Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 4A

The Little River Temperature (Fed ID: 41518) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013. The initial 2018 listing for exceedances of the Class VI (20°C) Natural Trout Waters criterion applies to 4.53 miles of the West Fork Little River. The West Fork Little River Aquatic Life Use impairment is nested in the Little River Temperature (Fed ID: 41518) TMDL Study.

West Fork Little River: Length 4.53 miles.

9-LWF004.55 (Rt. 8 Bridge North of Floyd) - The 2018 data window finds four of 12 temperature measurements exceeding the Class VI Natural Trout Waters 20°C criterion. Excursions are found on the following sampling dates and temperature measurements: 6/22/15 at 23.4°C, 7/23/15 at 21.2°C, 8/20/15 at 21.3°C, and 9/2/15 at 21.9°C.

		Cycle	TMDL	
	Cause	First	Dev.	Water
Assessment Unit / Water Name / Location Desc.	Category Cause Name	Listed	Priority	Size
VAW-N20R_LWF01A00 / West Fork Little River / West Fork Li	ittle 4A Temperature	2018	L	4.53
River waters from its mouth on Little River upstream to the mouth	of			

Dodd Creek on the West Fork Little River (NE51).

West Fork Little River Estuary Reservoir River (Sq. Miles) (Miles) (Acres) **Aquatic Life** 4.53

Temperature - Total Impaired Size by Water Type:

Sources:

Source Unknown

New River Basin

Cause Group Code: N20R-04-BEN Dodd Creek, Unnamed Tributary (XEM)

Cause Location: Unnamed tributary XEM from its mouth on Unnamed tributary XEL upstream to its headwaters (NE51).

City / County: Floyd Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 5A

This initial 0.71 mile 2018 data window Aquatic Life Use listing is based on Virginia Stream Condition Index scores collected as part of a special study.

9-XEM000.36 (Unnamed tributary (XEL) to Dodd Cr., UT)- Two 2016 VSCI scores define the Aquatic Life Use Impairment: Spring 39.0, Fall 63.9. This stream originates downslope of the Floyd County landfill (landfill was built on top of the original stream channel). Approximately 0.18 miles upstream of the sample station (9-XEM000.36), the stream surfaces from a spring box and is impacted by growths of iron bacteria and Sphaerotilus (sewage fungus). The spring 2016 sediment discharge appears to be affecting the benthic community. Certain stonefly taxa are tolerant of iron precipitate and can thrive in streams moderately impacted by landfills and mines.

9-XEM000.30 (Unnamed tributary (XEL) to Dodd Cr., UT) - Three VSCI spring surveys (2011-2012, 2016) report an average score of 58.2. The stream surfaces from the Floyd landfill in a spring box and is impacted by growths of iron bacteria and sphaerotilus (sewage fungus). The stream substrate was too impacted by bacterial growth to sample for benthic macroinvertebrates; A May 2011 habitat survey shows most parameters are in the optimal range. Sediment deposition was the only parameter found to be in the marginal range. During the 2012 habitat survey, scores for sediment deposition and several other parameters had declined, some were due to lower stream flow. The 2011 sample is dominated by mayflies, stoneflies and other generally pollution-sensitive taxa. The dominant mayfly taxa, Ephemerellidae (50% of all organisms) is somewhat tolerant of excessive sediment and several stonefly taxa present are known to be tolerant of iron precipitate and organic enrichment. The June 2012 sample finds the number of mayflies very low but stoneflies (51%) are numerous.

Assessment Unit / Water Name / Location Desc.	Cause Category Cau	se Name	Cycle First Listed	TMDL Dev. Priority	Water Size
/AW-N20R_XEM01A08 / Unnamed Tributary (XEM) / Unnam ributary XEM from its mouth on Unnamed tributary XEL upstread to the deadwaters (NE51).		Macroinvertebrates essments	2018	L	0.71
Dodd Creek, Unnamed Tributary (XEM)		Estuary		servoir	River
Dodd Creek, Unnamed Tributary (XEM) Aquatic Life		Estuary (Sq. Miles		servoir cres)	River (Miles)

Sources:

Upstream Source

New River Basin

Cause Group Code: N21R-01-BAC Little River (Lower)

Cause Location: The upper limit begins at the confluence of Dodd Creek (N19R) extending downstream to the Little River mouth on

the New River (N21R).

City / County: Floyd Co. Montgomery Co. Pulaski Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

Fecal Coliform / 4A

The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved 3/14/2012 and SWCB approved 3/25/2013. Exceedances of the former WQS fecal coliform (FC) bacteria instantaneous criterion of 1000 cfu/100 ml required the initial 2002 bacteria 303(d) Listing based on data from USGS station 03170000 where 2 of 14 observations exceed the criterion. Application of the revised 400 cfu/100 ml instantaneous criterion would result in 4 of 14 exceedances above the former criterion ranging from 420-14,900 cfu/100 ml. Due to the 2002 1.39 mi 303(d) Listing from Meadow Creek confluence downstream to the backwaters of Little River Reservoir and 2004 bacteria results from 9-LRV000.34 the impairment is extended 0.49 mi downstream. The 2012 Integrated Report (IR) extends the upper limit to the confluence of Dodd Creek incorporating the West Fork of Little River. The West Fork of Little River is nested within the overall Little River Bacteria TMDL. The impounded waters (60.44 ac) of Little River Reservoir are now bacteria impaired and were incorporated with the 2008 IR.

The 2004 IR establishes a 13.41 mi bacteria impairment at 9-LRV032.72 where 3/8 FC observations exceed the former WQS 400 cfu/100 ml instantaneous criterion. Exceedances range from 600-1100 cfu/100 ml. The impaired extent is from the end of Rt. 706 downstream to the confluence of Sidney Creek. This portion of Little River is separate from the original 2002 bacteria 303(d) Listing because of hydrology and the lack of bacteria data between the two initial listings on the mainstem of Little River.

Additional bacteria sample collection within the 2008 and 2010 data windows define the entire 44.22 mi impairment below. Escherichia coli (E.coli) bacteria replaces FC as the indicator per WQS [9 VAC 25-260-170. Bacteria; other waters].

West Fork Little River (Nested):

9-LWF004.55 (Rt 8 Br, N of Floyd) No additional data beyond the 2012 IR: 5/12 E.coli samples exceed the 235 cfu/100 ml instantaneous criterion with range from 280 to >2000 cfu/100 ml.

Little River

Final 2020

9-LRV044.49 (Rt. 615 Br) No additional data beyond the 2008 IR: 3/11 within the 2008 & 2010 data windows w/exceeding values from 380 to >2000 cfu/100 ml. 2006: 2/8 at 380 & 450 cfu/100 ml.

9-LRV032.72 (Rt. 617 Br) No additional data beyond the 2006 IR: 4/11 from 600-3300 cfu/100 ml. 2008: same as 2006 IR. 2010: remaining FC data = 1/3 samples in excess of the former instantaneous criterion at 3300 cfu/100 ml & no E.coli is available to assess.

9-LRV016.68 (Rt. 787 Br) 2018: 1/12 at 512 cfu/100 ml. 2012 & 2014: 2/12 at 380 & 1200 cfu/100 ml. 2010: 1/2 at >8000 cfu/100 ml. There are no E.coli data to assess. 2006 & 2008: 2/10 at 900 & > 8000 cfu/100 ml. There are no additional beyond the 2006 IR.

9-LRV012.58 (Rt. 787 pull off) 2016: 3/12 from 275-1075 cfu/100 ml.

9-LRV009.11 (Route 693 Bridge at Graysontown) 2018: 11/36; 2016: 10/36; 2014: 5/24 each cycle shows exceedances from 400 to >2000 cfu/100 ml. 2012: 2/12 at 400 & 1000 cfu/100 ml. 2010: 1/3 at 500 cfu/100 ml. 2006 & 2008 IRs: FC exceeds the former instantaneous criterion in 2/11 samples at 500 & 600 cfu/100 ml. Note: USGS 03170000 (Little R. at Graysontown) an original 2002 listing station is at the same location.

9-LRV000.44 (Above Little River Dam) No additional data beyond the 2010 IR: 2/7 at 420 & 1000 cfu/100 ml.

9-LRV000.34 (Rt. 605 Br- below Little River Dam) 2012 & 2010: 3/12 from 250 to >2000 cfu/100 ml. 2008 & 2006: 4/14 FC samples in excess of the former 400 cfu/100 ml criterion from 500-7300 cfu/100 ml. No additional data available beyond the 2012 IR.

Cause Category Cause Name

Cycle TMDL First Dev. Water Listed Priority Size

Assessment Unit / Water Name / Location Desc.

New River Basin							
	Fork Little River / West Fork Little ittle River upstream to the mouth of ttle River (NE51).	4A	Escherichia coli (E. coli))	2012	L	4.53
VAW-N21L_LRV01A02 / Little Reservoir from its impounding st	River Reservoir / Little River ructure upstream to its backwaters.	4A	Escherichia coli (E. coli)	2008	L	60.44
	River / The mainstem waters of New River upstream to the Little	4A	Escherichia coli (E. coli)	2010	L	0.49
	River / Mainstem Little River from servoir upstream to the end of the e Radford City intake (NE56).	4A	Escherichia coli (E. coli)	2012	L	0.69
	River / Mainstem Little River from am to the mouth of Meadow Creek	4A	Escherichia coli (E. coli))	2012	L	0.70
	River / The Little River mainstem w Creek upstream to the mouth of	4A	Escherichia coli (E. coli)	2012	L	12.33
	River / The Little River mainstem ek upstream to the WQS designated 8).	4A 1	Escherichia coli (E. coli)	2014	L	8.37
	River / Little River mainstem from at waters upstream to the mouth of 52).	4A	Escherichia coli (E. coli))	2006	L	3.70
Little River (Lower) Recreation				Estuary (Sq. Miles)		servoir cres)	River (Miles)
	Escherichia coli (E. coli) - Total Im	paire	d Size by Water Type:			60.44	30.81
Assessment Unit / Water N	ame / Location Desc. C	Cause Catego	e rry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
	River / Little River from the Brush confluence of Sidney Creek (NE53).	4A	Fecal Coliform		2004	L	8.79
	River / Little River from the end of	4A	Fecal Coliform		2004	L	4.62
Little River (Lower)				Estuary (Sq. Miles)		servoir .cres)	River (Miles)
	Fecal Coliform - Total Im	paire	d Size by Water Type:				13.41
Sources:							
Livestock (Grazing or Feeding Operations)		Unspe Waste	cified Domestic	Wet Wet (Non-P		Discharge urce)	es
Wildlife Other than							

New River Basin

Cause Group Code: N21R-02-BAC Meadow Creek

Cause Location: The Meadow Creek mainstem from the Mill Creek confluence downstream to the Meadow Creek mouth on Little

River (Radford South Quad).

City / County: Montgomery Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board

(SWCB) approved 3/25/2013.

Fecal coliform (FC) excursions of the former 1000 cfu/100 ml instantaneous criterion found in 2002 results in the initial 303(d) Listing of these waters for 4.49 miles. Exceedances are found in three of four observations and one geometric mean calculation exceedance is recorded in excess of the former 2002 criterion of 200 cfu/100 ml. Additional sample collections within the 2004 IR data window also produce exceedances of the former 400 cfu/100 ml instantaneous criterion in seven of 12 observations with one geometric mean excursion of the former criterion. Escherichia coli (E.coli) replaces fecal coliform (FC) bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters].

9-MDW004.62- There are no new data. The 2014, 2012 and 2010 data windows produce six exceeding values from 12 observations of the escherichia coli (E.coli) 235 cfu/100 ml instantaneous criterion. Values in excess of the criterion range from 280 to greater than 2000 cfu/100 ml. The 2006 IR finds FC exceedances of the former WQS 400 cfu/100 ml instantaneous criterion in six of 11 observations. The range of exceeding values is from 700 to greater than 8000 cfu/100 ml. FC exceedances and total observations within the 2008 data window are the same.

			Cycle	IMDL	
	Cause	•	First	Dev.	Water
Assessment Unit / Water Name / Location Desc.	Catego	ry Cause Name	Listed	Priority	Size
VAW-N21R_MDW01A00 / Meadow Creek / The Meadow Creek mainstem from its confluence with Little River upstream to the mou of Mill Creek on Meadow Creek (NE56).		Escherichia coli (E. coli)	2010	L	4.64

Meadow Creek		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			4.64

Sources:

Livestock (Grazing or Feeding Operations)

On-site Treatment Systems

Unspecified Domestic Wet Weather Discharges
(Septic Systems and Similar Waste
Decentralized Systems)

Wet Weather Discharges
(Non-Point Source)

Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N21R-03-BAC Mill Creek, Poplar Branch, Mill Creek UTs (XDE & XDF)

Cause Location: The upper limit begins at the headwaters of Mill Creek on the Riner Quad and extends downstream to the Mill

Creek confluence with Meadow Creek at the Rt. 600 Bridge on the Radford South Quad (7.04 miles). This impairment also includes Poplar Branch and its tributaries form its mouth on Mill Creek to its headwaters as well as

to unnamed tributaries to Mill Creek (XDE & XDF).

City / County: Montgomery Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

Fecal Coliform / 4A

The Mill Creek Bacteria Total Maximum Daily Load (TMDL) is U.S. EPA approved 6/5/2002 [Fed ID 9453/19986] and SWCB approved 6/17/2004 (formerly VAW-N21R-03). The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved 3/14/2012 and SWCB approved 3/25/2013. The Mill Creek Bacteria Implementation Plan (IP) received SWCB approval 6/27/2007; Little River 3/25/2013. The 1996/2002/2004 impaired waters now extend to the headwaters of Mill Creek (7.60 mi). 2002 tributary additions include Poplar Branch and 2 unnamed tributaries. The waters are impaired for a total of 15.92 mi.

The waters are originally 303(d) Listed based on the former fecal coliform (FC) WQS instantaneous criterion of 1000 cfu/100 ml and 200 geometric mean. The 2004 Integrated Report (IR) records exceedances of both the former FC 400 cfu/100 ml instantaneous criterion and geometric mean criterion of 200 cfu/100 ml. Listed below are the monitored sites showing FC instantaneous excursions / total sample collections; (maximum) and geomean calculation exceedances / total calculations, where applicable. Instantaneous Escherichia coli (E. coli) single observations from the 2008 IR are listed next (value). Each exceed the WQS instantaneous criterion of 235 cfu/100 ml. Escherichia coli (E.coli) replaces FC bacteria as the indicator as per WQS [9 VAC 25-260-170. Bacteria; other waters].

9-MLC005.44- There are no new data since the 2014 IR: 18/36 from 250 to >2000 cfu/100 ml. 2012: 11/24 from 280-600 cfu/100 ml. 2010: 4/12 from 250-580 cfu/100 ml.

9-MLC002.59 (Rt. 669 Bridge) There are no new data since the 2014 IR: 23/36. 2014: 16/24 from 280 to >2000 cfu/100 ml. 2012: 14/24 from 280 to >2000 cfu/100 ml. 2010: 7/12 from 580 to >2000 cfu/100 ml.

9-MLC001.53 (Rt. 693, Childress) There are no new data since the 2014 IR: 14/36 from 250 to >2000 cfu/100 ml. 2012: 8/24 from 250-1100 cfu/100 ml. 2010: 3/12 from 300-1100 cfu/100 ml.

Data below reflect the 2004, 2006 and 2008 data windows as there were no additional data beyond the 2006 IR. Two ambient fixed sites 9-MLC005.44 & 9-MLC001.53 are included with the non-fixed sites below.

2004 IR results:

Mill Creek

9-MLC000.17 (Rt. 600 Bridge) - 3/5; (3900); 1/1 geomean; E.coli- 1/1 (800).

9-MLC001.31 (Rt. 693 Bridge) - 3/5; (2300); 1/1 geomean; E.coli- 1/1 (800).

9-MLC001.53 (Rt. 693, Childress) - 3/6; (2300).

9-MLC002.74 (Private Road off Rt. 616) - 4/5; (>8000); 1/1 geomean; E.coli- 1/1 (800).

9-MLC005.44 (Rt. 8 Bridge-above Riner STP) - 18/25; (2500); E.coli- 1/1 (800).

9-MLC006.00 (Private road Rt. 616) - 2/5; (>8000); 0/1 geomean; E.coli- 1/1 (>800).

Poplar Branch

9-PPL000.01 (Private Road at mouth) - 1/1; (>8000).

9-PPL001.27 (Rt. 616 Bridge) - 2/2 (2800).

Mill Creek Unnamed Tributaries

9-XDE000.95 (Rt. 678 Bridge) - 4/5; (>8000); 1/1 geomean; E.coli- 1/1 (>800).

9-XDF000.11 (Private road Rt. 669) - 4/5;(2600); 1/1 geomean; E.coli- 1/1 (>800).

2006 IR results for 2006 stations within the data window:

Mill Creek

9-MLC005.44- 2006 FC exceeds the 400 cfu/100 ml instantaneous criterion in 10/15 obs ranging from 600-2000 cfu/100 ml. 2008: 8/11.

9-MLC002.74- 2006 FC exceeds in 10/12 obs (max. > 8000 and min. 500 cfu/100 ml). 2008: 9/11.

New River Basin

Wildlife Other than Waterfowl

9-MLC001.53- 2006 FC exceeds in 5/8 (max. 2300 cfu/100 ml). 2008: 5/8.

Assessment Unit / Water	Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
	Creek / Mill Creek mainstem waters ek upstream to the Montgomery (NE56).	s 4A	Escherichia coli (E. coli)	2010	L	5.49
	Creek / Mill Creek mainstem waters SA Riner STP outfall upstream to its		Escherichia coli (E. coli)	2010	L	2.11
Mill Creek, Poplar Branch, M	III Creek UTs (XDE & XDF)			Estuary	Res	servoir	River
Recreation				(Sq. Miles)) (A	cres)	(Miles)
	Escherichia coli (E. coli) - Total I	mpaired	Size by Water Type:				7.60
Assessment Unit / Water	Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
	lar Branch & Tributaries / Poplar es from its confluence with Mill Creek E56).		Fecal Coliform		2002	L	4.62
	Creek, UT (XDE) / An unnamed om its mouth upstream. The stream i ill Creek flowing to VAW-	4A is	Fecal Coliform		2002	L	1.75
	Creek, UT (XDF) / An unnamed om its mouth upstream. The stream i ill Creek flowing to VAW-	4A	Fecal Coliform		2002	L	1.95
Mill Creek, Poplar Branch, M	III Creek UTs (XDE & XDF)			Estuary (Sq. Miles)		servoir .cres)	River (Miles)
	Fecal Coliform - Total I	mpaired	Size by Water Type:				8.32
Sources:							
Livestock (Grazing or Feeding Operations)	On-site Treatment Systems (Septic Systems and Similar Decentralized Systems)	Unspe Waste	cified Domestic		eather loint So	Dischargource)	es

New River Basin

Cause Group Code: N21R-05-BAC **Brush Creek**

Cause Location: Brush Creek from the first bridge on Route 617 south of the junction of Routes 617 and 601 downstream to the

Brush Creek mouth on Little River.

City / County: Montgomery Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013.

The 2004 Recreational Use impairment continues for 5.94 miles originally due to fecal coliform (FC) bacteria exceedances of the former instantaneous criterion of 400 cfu/100 ml. Escherichia coli (E.coli) replaces fecal coliform (FC) bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters].

9-BSH000.05 (Rt. 617 Bridge) Four of 12 E.coli samples exceed the 235 cfu/100 ml instantaneous criterion within the 2018 data window. Excursions range from 473 to greater than 1100 cfu/100 ml. The 2012 Integrated Report (IR) found six of 11 escherichia coli (E.coli) observations exceed the 235 cfu/100 ml instantaneous criterion. Excessive values range from 250 to greater than 2000 cfu/100 ml. These waters were initially Listed for fecal coliform (FC) in 2004 with three of eight FC samples exceeding the former WQS instantaneous criterion of 400 cfu/10 ml. The 2010 data window finds two of two samples exceeding the former instantaneous criterion at 800 and 1100 cfu/100 ml. The 2004, 2006 and 2008 data windows find five of 10 FC samples exceeding the former instantaneous criterion. The maximum exceedance range is from 700 to 1300 cfu/100 ml. There were no E.coli data to assess at that time.

			Cycle	TMDL	
	Cause	Э	First	Dev.	Water
Assessment Unit / Water Name / Location Desc.	Catego	ry Cause Name	Listed	Priority	Size
VAW-N21R_BSH01A04 / Brush Creek / Brush Creek from the	first 4A	Escherichia coli (E. coli)	2012	L	5.94
bridge on Route 617 south of the junction of Routes 617 and 601					
downstream to the Brush Creek mouth on Little River (NE52).					

Brush Creek		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type) :		5.94

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

Sources:

Livestock (Grazing or **On-site Treatment Systems Unspecified Domestic** Wet Weather Discharges Feeding Operations) (Septic Systems and Similar Waste (Non-Point Source)

Decentralized Systems)

Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N21R-06-BAC **Laurel Creek**

Cause Location: Laurel Creek mainstem from its headwaters NW of the Routes 608 and 673 intersection downstream to its

confluence with Little River.

City / County: Floyd Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board

(SWCB) approved 3/25/2013.

Fecal coliform (FC) bacteria exceedances cause this initial 2004 303(d) Listed water to not support the Recreational Use for 3.44 miles. Escherichia coli (E.coli) replaces fecal coliform (FC) bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters].

9-LLL000.05- There are no additional data beyond the 2012 Integrated Report (IR) where five of 12 escherichia coli (E.coli) samples exceed the 235 cfu/100 ml instantaneous criterion. The range of exceeding values is 380 to 1200 cfu/10 ml. FC data within the 2010 data window find one of two samples in excess of the former criterion. The single exceedance is 1000 cfu/100 ml. Both the 2006 and 2008 Integrated Reports (IR) find FC exceeds the former WQS 400 cfu/100 ml instantaneous criterion in five of 10 samples. The exceedances range from 600 to 2800 cfu/100 ml. FC data within the 2010 data window find one of two samples in excess of the former criterion. The single exceedance is 1000 cfu/100 ml.

Cycle TMDI

Sources:	On alta Transfer and Outland	Harris de Demontis	\\/ a + \\/ a	- 41 -	N: b	
	Escherichia coli (E. coli) - Tota	al Impaired Size by Water Type:				3.44
Laurel Creek Recreation			Estuary (Sq. Miles)		ervoir cres)	River (Miles)
VAW-N21R_LLL01A04 / Lau neadwaters (Class VI) NW of I downstream to its confluence		4A Escherichia coli (E. coli	2	2012	L	3.44
Assessment Unit / Water	Name / Location Desc.	Cause Category Cause Name	j	First isted	Dev. Priority	Water Size

Livestock (Grazing or **On-site Treatment Systems Unspecified Domestic** Wet Weather Discharges (Septic Systems and Similar Feeding Operations) Waste (Non-Point Source) Decentralized Systems)

Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N21R-07-BAC Big Indian Creek

Cause Location: Big Indian Creek from approximately 0.5 miles upstream of the West Fork Big Indian Creek mouth downstream to

the Big Indian Creek confluence with Little River.

City / County: Floyd Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013. The 2010 Integrated Report initially 303(d) Lists these waters.

9-BIC000.14 (Rt. 787 Bridge)- There is no additional bacteria data collected within the 2018 or 2016 data windows. 2014 data window reveals 10 of 24 escherichia coli (E.coli) observations in excess of the 235 cfu/100 ml instantaneous criterion. Excursions range from 300 to 950 cfu/100 ml. There are no additional data within the 2012 data window. The 2010 initial Listing is based on E.coli exceedances from four of 12 samples in excess of the instantaneous criterion with excursions ranging from 350 to 950 cfu/100 ml.

Cuala TMDI

		Cycle			
	Cause	First	Dev.	Water	
Assessment Unit / Water Name / Location Desc.	Category Cause Name	Listed	Priority	Size	
/AW-N21R_BIC01A02 / Big Indian Creek / Big Indian Creek	4A Escherichia coli (E. coli)	2010	L	7.83	

VAW-N21R_BIC01A02 / Big Indian Creek / Big Indian Creek mainstem from approximately 0.5 miles upstream of the West Fork Big Indian Creek mouth downstream to the Big Indian Creek confluence with Little River (NE54).

Big Indian Creek		Estuary	Reservoir	River
Recreation		(Sq. Miles)	(Acres)	(Miles)
	Escherichia coli (E. coli) - Total Impaired Size by Water Type:			7.83

Sources:

Livestock (Grazing or Feeding Operations) On-site Treatment Systems Unspr (Septic Systems and Similar Waster Decentralized Systems)	ecified Domestic Wet Weather Discharges (Non-Point Source)
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Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N21R-07-TEMP **Big Indian Creek**

Cause Location: Big Indian Creek from approximately 0.5 miles upstream of the West Fork Big Indian Creek mouth downstream to

the Big Indian Creek confluence with Little River.

City / County: Floyd Co. Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 4A

The Little River Temperature (Fed ID: 41518) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved on 3/25/2013. These waters were initially 303(d) Listed with the 2004 assessment and subsequently delisted with the 2010 assessment. The waters return to an impaired status with the 2014 assessment. Big Indian Creek is addressed by the Little River Temperature TMDL and is category 4A.

9-BIC000.14- (Rt. 787 Bridge, Indian Valley Rd.) No new data was collected within the 2018 data window. The 2016 data windows finds no new temperature exceedances. The 2014 data window records three of 24 temperature measurements in excess of the Class V stockable trout water criterion of 21°C. Exceedances occur on 7/18/2007 at 21.7°C; 7/21/2011 at 24.1°C and 8/29/2012 at 21.6°C. The 2012 data window reveals one exceeding value at 21.7°C on 7/18/2007 from 12 measurements with no additional data. The waters were delisted based on data within the 2010 window where one exceedance (7/18/2007) is recorded from 15 measurements. The temperature original 2004 303(d) Listing continued through the 2008 Cycle. 2006 and 2008 IRs record two of 11 exceedances each. The excursions are 23.9 °C on 7/11/01 and 23.2 °C on 7/10/02 during some of the driest years on record. The original 303(d) Listing in 2004 is based on two of eight temperature measurements exceeding the 21°C criterion as recorded for 2006 and 2008 data windows.

	Cause	First	Dev.	Water
Assessment Unit / Water Name / Location Desc.	Category Cause Name	Listed	Priority	Size
VAW-N21R_BIC01A02 / Big Indian Creek / Big Indian Creek	4A Temperature	2004	L	7.83
mainstem from approximately 0.5 miles upstream of the West For	k			
Big Indian Creek mouth downstream to the Big Indian Creek				

confluence with Little River (NE54).

Big Indian Creek		Estuary	Reservoir	River
Aquatic Life		(Sq. Miles)	(Acres)	(Miles)
	Temperature - Total Impaired Size by Water Type:			7.83

Sources:

Natural Sources

New River Basin

Cause Group Code: N21R-08-BAC Beaver Creek

Cause Location: Beaver Creek from its mouth on Little River to its headwaters.

City / County: Floyd Co.
Use(s): Recreation

Wildlife Other than Waterfowl

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The Little River Bacteria (Fed ID: 41519) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved 3/25/2013. The 2020 Integrated Report initially 303(d) Lists these waters. Beaver Creek is Nested in the aforementioned TMDL Study.

9-BVR001.84 (Rt. 705 Bridge, Floyd Co.) - The 2020 data window finds seven of 12 Escherichia coli (E.coli) samples in exceedance of the 235 cfu/100 ml instantaneous criterion. The minimum excursion was 327 cfu/100 ml (9/25/18) and maximum 2,481 cfu/100 ml (6/21/18).

Assessment Unit / V	Vater Name / Location Desc.	Cause Category Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N21R_BVR01A20 mouth on Little River to it	/ Beaver Creek / Beaver Creek from its s headwaters (NE52).	4A Escherichia coli (E. coli)		2020	L	7.11
Beaver Creek Recreation			Estuary (Sq. Miles)		servoir acres)	River (Miles)
	Escherichia coli (E. coli) - Total	Impaired Size by Water Type:				7.11
Sources:						
Livestock (Grazing o Feeding Operations)	On-site Treatment Systems (Septic Systems and Similar Decentralized Systems)	Unspecified Domestic Waste	Wastes	s from F	Pets	

New River Basin

Cause Group Code: N21R-08-BEN Beaver Creek

Cause Location: Beaver Creek from its mouth on Little River to its headwaters.

City / County: Floyd Co. Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

The Little River Sediment (Fed ID: 41517) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved on 3/25/2013. These waters are initially 303(d) Listed with the 2020 assessment. Beaver Creek is nested in the Little River Sediment TMDL and is category 4A.

9-BVR001.70 (Off Rt. 705 / Beaver Cr. Rd.) - Bio 'IM' from two 2018 VSCI Scores of 48.3 (S) and 58.3 (F). This station was surveyed as part of the Probabilistic monitoring program in 2018. The benthic macroinvertebrate community had good diversity but was comprised of an even mix of both pollution tolerant and sensitive taxa.

Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
$VAW-N21R_BVR01A20\ /\ Beaver\ Creek\ /\ Beaver\ Creek\ from\ its\ mouth\ on\ Little\ River\ to\ its\ headwaters\ (NE52).$	4A Benthic Macroinvertebrat Bioassessments	tes	2020	L	7.11
Beaver Creek		Estuary (Sq. Miles)		servoir cres)	River (Miles)
Aquatic Life Benthic Macroinvertebrates Bioassessments - Total	Impaired Size by Water Type:	(Oq. IVIIICS)	(/\	0103)	7.11

Sources:

Erosion from Derelict Land Livestock (Grazing or Barren Land) Loss of Riparian Habitat Non-Point Source (Barren Land) Feeding Operations)

Sediment Resuspension Streambank

(Clean Sediment) Modifications/Destabilization

New River Basin

Cause Group Code: N21R-08-TEMP **Beaver Creek**

Cause Location: Beaver Creek from its mouth on Little River to its headwaters.

City / County: Floyd Co. Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 4A

The Little River Temperature (Fed ID: 41518) TMDL Study is U.S. EPA approved on 3/14/2012 and State Water Control Board (SWCB) approved on 3/25/2013. These waters are initially 303(d) Listed with the 2020 assessment and are nested in the Little River Temperature TMDL (Category 4A).

9-BVR001.84 (Rt. 705 Bridge, Floyd Co.) - The 2020 data window finds three of 12 temperature measurements in exceedance of the 20 C Class VI Natural Trout Waters Criteria. Excursions are: 21 C (6/21/18), 21 C (7/23/18), 22 C (8/13/18).

			Cycle	IMDL		
	Cause	e	First	Dev.	Water	
Assessment Unit / Water Name / Location Desc.	Catego	ry Cause Name	Listed	Priority	Size	
VAW-N21R_BVR01A20 / Beaver Creek / Beaver Creek from its	4A	Temperature	2020	L	7.11	
mouth on Little River to its headwaters (NE52).						

Beaver Creek Estuary Reservoir River (Sq. Miles) (Acres) (Miles) **Aquatic Life** 7.11

Temperature - Total Impaired Size by Water Type:

Sources:

Loss of Riparian Habitat Streambank

Modifications/Destabilization

New River Basin

Cause Group Code: N22R-02-BAC Stroubles Creek

Cause Location: The upstream end is at the Duck Pond dam on the southwest end of the VPI&SU campus on the Blacksburg Quad.

The downstream end is at the Slate Branch mouth on Stroubles Creek.

City / County: Montgomery Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

Fecal coliform (FC) bacteria exceedances of the former 1000 cfu/100 ml WQS instantaneous criterion in 2002 cause impairment of the Recreational Use. Three of 23 observations exceed the former criterion at station 9-STE002.41 Rt. 705 Bridge (Coal Hollow Road). The 2004 IR at 9-STE002.41 records four exceedances from 35 samples in excess of the current 400 cfu/100 ml WQS instantaneous criterion. Escherichia coli (E.coli) bacteria replaced fecal coliform (FC) in 2006 as the indicator as required by Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters]. 2008 results find E.coli exceedances at 9-STE002.41 are three of 31 samples and resulted in 2.11 miles delisted with the 2008 IR. This 2.11 mile delisted portion (partial - length) returned with the 2010 303(d) Listing.

9-STE002.41- (Rt. 705 Bridge- Coal Hollow Road) Nine of 36 excursions are reported during the 2020 data window. The 2018 and 2016 data windows find eleven of 36 escherichia coli (E.coli) samples that exceed the 235 cfu/100 ml instantaneous criterion. Twelve of 36 samples exceeded that criterion in 2014. Values in excess of the instantaneous criterion range from 250 to greater than 2000 cfu/100 ml within the 2014 and 2016 data windows. The 2012 data window finds eight of 36 observations exceeding the 235 cfu/100 ml instantaneous criterion. Exceeding values range from 280 to greater than 2000 cfu/100 ml. 2010 E.coli samples find eight exceed the 235 cfu/100 ml instantaneous criterion from a total of 32 samples with the same range of exceedance.

9-STE007.29 (Rt. 657 Bridge below old B'Burg STP) E.coli exceeds in seven of 11 during the 2020 data window. There is no new data within the 2016 data window. Escherichia coli (E. coli) samples find eight exceed the 235 cfu/100 ml instantaneous criterion from a total of 24 samples. Exceeding values range from 280 to greater than 2000 cfu/100 ml within the 2014 data window. The 2012 IR reports eight E.coli samples exceed the instantaneous criterion from a total of 33. Exceeding values range from 280 to greater than 2000 cfu/100 ml. 2010 results find eight exceed from a total of 32 samples with the same range of exceedance as 2012. 2008 E.coli results exceed in five of 25 samples. The 2008 exceedance range is from 300 to greater than 2000 cfu/100 ml. 2006 E.coli samples reveal five exceed the instantaneous criterion from a total of 16. Exceeding values range from 490 to greater than 5000 cfu/100 ml.

Assessment Unit / Water Na	ime / Location Desc.	Cause Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N22R_STE03A00 / Stroub waters extend from the Slate Bran upstream to the mouth of Walls B	nch mouth on Stroubles Creek	5A	Escherichia coli (E. coli)		2010	Н	2.11
VAW-N22R_STE04A00 / Stroub waters extend from the Walls Bra Pond located on the VPI&SU Car	nch mouth upstream to the Duck	5A	Escherichia coli (E. coli)		2006	Н	5.08
Stroubles Creek Recreation				Estuary (Sq. Miles)		servoir cres)	River (Miles)
	Escherichia coli (E. coli) - Total	Impaire	d Size by Water Type:				7.19
Sources:							
Discharges from Municipal Separate Storm Sewer Systems (MS4)	Livestock (Grazing or Feeding Operations)	Municipal (Urbanized High Density Area)		Unspec Waste	cified Do	omestic	
Wastes from Pets	Wet Weather Discharges (Non-Point Source)	Wildlif Water	e Other than fowl				

New River Basin

Cause Group Code: N22R-02-BEN Stroubles Creek

Cause Location: These mainstem waters extend from the Walls Branch mouth upstream to the Duck Pond located on the VPI&SU

Campus.

City / County: Montgomery Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

The Stroubles Creek General Standard (Benthic- Sediment) Total Maximum Daily Load (TMDL) is U.S. EPA approved on 1/28/2004 [Fed ID: 21904]. The SWCB approved the Study on 6/17/2004. The Benthic (Sediment) Implementation Plan (IP) is SWCB approved (9/27/2006) (formerly VAW-N22R-02). The 1996 original 303(d) Listed waters remain impaired for contravention of the General Standard (Benthic).

9-STE007.29 - (Rt. 657 Bridge below old Blacksburg STP) Bio 'IM' from six VSCI scores (2015-17) averaging 57.9. The 2018 data window includes six VSCI scores averaging 54.6 (2012, 2015-2016). The Spring 2016 score indicated improvement from Spring 2015 and the Fall 2016 score maintained a Non-Impaired status. While overall the VSCI scores indicate an impaired community, the scores improved during this assessment period. The 2014 assessment found nine Virginia Stream Condition Index (VSCI) surveys (2007-2010 & 2012) are 'IM' with an average score of 46.82. Impairment is found from nine surveys (2006 - 2010) with an average score of 46.82 in 2012. The 2010 assessment found impairment from seven VSCI surveys (2003 & 2006 - 2008) with an average score of 45.6. An average score of 45.6 is also found in 2008 from six VSCI surveys (2001 - 2003 & 2006).

The moderately pollution tolerant caddisfly family Hydropsychidae and fly family Chironomidae were the second most common macroinvertebrates during these surveys. This community indicates the benthic community is exposed to moderate level of pollution, possibly a nutrient source that provides the Hydropsychidae the opportunity to be second most dominant. Thus, this stream reach shows evidence of year long pollution. Habitat condition at this station is suboptimal, impacted by sediment and poor riparian vegetation zones. The mostly open canopy allows for increased water temperatures and primary production resulting in large mats of algae and bacteria on the stream substrate during the summer and fall.

		Cycle	TMDL	
	Cause	First	Dev.	Water
Assessment Unit / Water Name / Location Desc.	Category Cause Name	Listed	Priority	Size

VAW-N22R_STE04A00 / Stroubles Creek / These mainstem waters extend from the Walls Branch mouth upstream to the Duck Pond located on the VPI&SU Campus (NE59).

Estuary Reservoir River (Sq. Miles) (Acres) (Miles)

1996

L

5.08

Benthic Macroinvertebrates

Bioassessments

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type: 5.08

Sources:

Stroubles Creek

Aquatic Life

Discharges from Municipal Livestock (Grazing or Separate Storm Sewer Feeding Operations)

Systems (MS4)

Livestock (Grazing or Municipal (Urbanized High Density Area)

Sediment Resuspension (Clean Sediment)

New River Basin

Cause Group Code: N22R-03-BAC Back Creek

Cause Location: The waters extend from 0.70 miles below the Rt. 636 Bridge crossing downstream to Back Creek's mouth on the

New River.

City / County: Pulaski Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The 1996 303(d) Listed Back Creek Bacteria Total Maximum Daily Load (TMDL) is U.S. EPA approved on 6/21/2004 [Fed ID 24564] and SWCB approval on 12/02/2004. The Bacteria/Benthic Implementation Plan (IP) is SWCB approved 7/31/2008 (formerly VAW-N22R-03). 1996 fecal coliform (FC) exceedances are found in seven of seven observations at 9-BCK009.47; 2002 records 17 of 23 samples exceeding the former fecal coliform bacteria instantaneous criterion of 1000 cfu/100 ml. The 2004 Integrated Report (IR) records 19 of 21 samples exceeding the former WQS fecal coliform bacteria instantaneous criterion of 400 cfu/100 ml at 9-BCK009.47. The excursions range from 900 to >8000 cfu/100 ml. Escherichia coli (E.coli) bacteria replaced fecal coliform in 2006 as the indicator as required by Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters]. The waters remain impaired for 17.53 miles with the 2014 and 2016 Assessments.

9-BCK015.98 (Rt. 636 Bridge, Black Hollow Road) Nine of 11 excursions reported at 9-BCK015.88 during the 2020 data window. Six of seven E.coli samples exceed the 235 cfu/100 ml instantaneous criterion within the 2018 data window. Prior to 2018, there are no additional data beyond the 2012 assessment where escherichia coli (E.coli) samples exceeded the WQS instantaneous criterion of 235 cfu/100 ml in 24 of 36 total samples. Values in excess of the criterion range from 300 to greater than 2000 cfu/100 ml. 2010 E.coli samples exceed the instantaneous criterion in 25 of 35 samples. Values in excess of the criterion range from 300 to greater than 2000 cfu/100 ml. The 2008 assessment finds E.coli exceeds the instantaneous criterion in 19 of 26 samples. Values in excess of the criterion range from 250 to greater than 2000 cfu/100 ml. In 2006 E.coli samples exceed the instantaneous criterion in 11 of 14 samples with the same exceedance range.

9-BCK009.47 (Rt. 100 Bridge) There are no additional data beyond the 2012 Integrated Report (IR) where E.coli exceeds the 235 cfu/100 ml criterion in 34 of 36 samples. The range of exceedance is from 320 to greater than 2000 cfu/100 ml. 2010 E. coli exceedances of the instantaneous criterion are found in 39 of 42 samples. The range of exceeding values is from 310 cfu/100 ml to 18,000. E.coli exceeds the instantaneous criterion in 32 of 35 samples in 2008. The range of exceeding values is from 310 cfu/100 ml to 18,000. Two of two geometric mean calculations exceed the 126 cfu/100 ml criterion based on the former WQS frequency of collection. The 2006 assessment found E.coli exceeds the instantaneous criterion in 20 of 21 samples with the same exceedance range.

9-BCK000.74 (Rt. 600 Bridge) There are no additional data beyond the 2012 IR where 20 of 36 E.coli exceedances occur ranging from 250 to greater than 2000 cfu/100 ml. E.coli exceedances are found in 29 of 43 samples within the 2010 data window. Exceedances range from 250 cfu/100 ml to 9000. The 2008 assessment finds E.coli exceeds the instantaneous criterion in 23 of 36 samples with exceedances ranging from 290 cfu/100 ml to greater than 2000. Three of three geometric mean calculations exceed the 126 cfu/100 ml criterion based on the former WQS frequency of collection. The exceedance range in 2006 is the same as 2008 where E.coli exceeds in 15 of 22 samples.

Cycle TMDI

Escherichia coli (E. coli) - Total Impaired Size by Water Type:					
Back Creek Recreation			eservoir Acres)	River (Miles)	
AW-N22R_BCK02A08 / Back Creek / Back Creek from 0.70 niles downstream of the Rt. 636 crossing on downstream to the onfluence of Shuffle Branch (NE61).	4A Escherichia coli (E. coli)	2006	L	11.77	
AW-N22R_BCK01A00 / Back Creek / Back Creek mainstem raters from the mouth of Shuffle Branch downstream to Back creek's mouth on the New River (NE61).	4A Escherichia coli (E. coli)	2006	L	5.76	
Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	First Listed		Water Size	

New River Basin

Sources:

Livestock (Grazing or Feeding Operations)

(Non-Point Source)

Wet Weather Discharges

On-site Treatment Systems (Septic Systems and Similar

Decentralized Systems)

Wildlife Other than Waterfowl

Unspecified Domestic Waste

Wastes from Pets

New River Basin

Cause Group Code: N22R-03-BEN Back Creek

Cause Location: The waters extend from 0.70 miles below the Rt. 636 Bridge crossing downstream to Back Creek's mouth on the

New River.

City / County: Pulaski Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

The 2002 303(d) Listed Back Creek General Standard (Benthic- Sediment) Total Maximum Daily Load (TMDL) is U.S. EPA approved on 6/21/2004 [Fed ID 24565]. The SWCB approved the Study on 12/02/2004. The Benthic/Bacteria Implementation Plan (IP) is SWCB approved 7/31/2008.

The TMDL identifies sediment as the primary stressor for the aquatic life use (benthic) impairment. The 2002 severe RBP II score of 37.50 produces the initial 17.53 mile listing of the benthic impairment. The 2008 assessment finds via station 9-BCK000.74 that a single Virginia Stream Condition Index (VSCI) score indicates full support. A potential delisting could occur for the lower end of Back Creek should additional surveys produce scores at 60 or above in succeeding assessment cycles.

9-BCK015.98- (Rt. 636 Bridge, Black Hollow Road) The 2018 data window finds Bio 'IM' from six VSCI scores. The addition of 2016 spring and fall data results in a VSCI averaging 44.7. Bio 'IM' Four Virginia Stream Condition Index (VSCI) surveys (2011-2012) within the 2016 and 2014 data windows produce an average score of 42.9. And two 2006 VSCI surveys with an average score of 42.8 are reported within previous Integrated Reports (IR). The habitat surveys indicate the stream is impacted by sediment deposition, riparian vegetation removal, channel alteration (straightening of the stream), and destabilized stream banks. Additionally, the water in Back Creek is often turbid from cattle disturbance of stream banks and in-stream sediments. These impacts result in stream substrate and water that limits colonization of benthic macroinvertebrates and fish.

9-BCK009.47 (Rt. 100 Bridge) Bio- 'IM'; The 2012 Integrated Report (IR) reveals four VSCI surveys (2006 & 2010) with an average score of 41.0 The remaining two surveys within the 2014 and 2016 data windows produce an average score of 32.6. The benthic community is dominated by taxa that are tolerant of nutrient/organic enrichment. Late summer of 2006 a fish kill occurred that was the probable cause for the decline in the benthic community for the Fall sample. The community recovered between Fall of 2006 and Spring of 2010, however a decline is noted in the Fall 2010 score. NPS pollution from agricultural sources upstream from Rt. 100 has impacted the stream. Habitat at this site has been impacted by the agricultural land use in the watershed, resulting in sedimentation and excessive algal growth on the substrate. The 2008 and 2010 assessments report three VSCI surveys (2003 & 2006) with an average score of 41.0 as well.

9-BCK000.74- (Rt. 600 Bridge) Bio- 'FS' There are no additional surveys beyond the 2006 IR. One fall 2003 VSCI survey scoring 67.2. This AU would be a candidate for delisting should additional surveys find scores above 60. The reach appears to have habitat that would suit a diverse benthic community and was surveyed to determine if it was a recovery zone from upstream impairments. However, this station had a low abundance of sensitive EPTs. The high dominance of Elmidae (53.3%) is possibly due to slight nutrient enrichment and the subsequent abundance of periphyton, which is the predominant food of riffle beetles. This station is slightly impacted by sediment deposition. The banks and riparian zones are impacted by altered hydrology and human activities. However, the substrate size, frequency of riffles, flow, velocity, and channel gradient have a positive effect on the benthic community.

Cvcle TMDL

Assessment Unit / Water Name / Location Desc.	Cause Catego	e rry Cause Name		First Listed	Dev. Priority	Water Size
VAW-N22R_BCK01A00 / Back Creek / Back Creek mainstem waters from the mouth of Shuffle Branch downstream to Back Creek's mouth on the New River (NE61).	4A	Benthic Macroinvertebrates Bioassessments	3	2002	L	5.76
VAW-N22R_BCK02A08 / Back Creek / Back Creek from 0.70 miles downstream of the Rt. 636 crossing on downstream to the confluence of Shuffle Branch (NE61).	4A	Benthic Macroinvertebrates Bioassessments	3	2002	L	11.77
Back Creek			Estuary Sg. Miles)		servoir cres)	River (Miles)
Aquatic Life Benthic Macroinvertebrates Bioassessments - Tota	l Impaire	`	,q. mico,	(7)	10100)	17.53

New River Basin

Sources:

Channelization

Loss of Riparian Habitat

Sediment Resuspension (Clean Sediment)

New River Basin

Cause Group Code: N22R-04-BAC Toms Creek

Cause Location: Toms Creek mainstem waters just below the Poverty Creek confluence upstream to its headwaters.

City / County: Montgomery Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

This initial 2014 Listing is a result of bacteria data showing impairment of the Recreational Use. The 2020 data window extends this impairment downstream 5.71 miles due to E.coli data collections at 9-TOM005.32.

9-TOM012.78- (Lower bike path off Deerfield Drive) Three of 12 escherichia coli (E.coli) samples exceed the 235 cfu/100 ml instantaneous criterion within the 2014 data window. Exceeding values range from 275 to 950 cfu/100 ml. Note: Level 2 Citizen data indicates the impairment extends downstream to the Toms Creek confluence with the New River. There was no additional data collected since the 2014 data window.

9-TOM005.32 (Rt. 725 Bridge [Poverty Cr Road]) - E.coli 'IM' from five excursions of the 235 cfu/100 ml instantaneous criterion from 12 samples during the 2020 data window.

Assessment Unit / Water N	Name / Location Desc.	Cause Category Cause Name	e	Cycle First Listed	TMDL Dev. Priority	Water Size
waters just below the Poverty C	as Creek / Toms Creek mainstem reek confluence upstream to the s are not within the WQS public wa 0).	5A Escherichia coli ater	(E. coli)	2020	L	5.71
VAW-N22R_TOM03A08 / Tommouth of Big Run upstream to it	s Creek / Toms Creek from the s headwaters (NE60).	5A Escherichia coli	(E. coli)	2014	Н	6.13
Toms Creek Recreation			Estuary (Sq. Miles		servoir .cres)	River (Miles)
	Escherichia coli (E. coli) - Total	Impaired Size by Water	Type:			11.84
Sources:						
Livestock (Grazing or Feeding Operations)	Municipal (Urbanized High Density Area)	Non-Point Source	(Septi	c Systen	nent Syst ns and Si Systems	milar
			20001	i.i.a.ioa	0,0000	,

New River Basin

Cause Group Code: N22R-04-TEMP Toms Creek

Cause Location: Toms Creek mainstem waters from its mouth on the New River upstream to its headwaters.

City / County: Montgomery Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5C

The initial 2008 5.71 mile impairment is extends upstream 6.13 miles (2012) and downstream 4.56 miles (2014) with data provided by the National Committee for the New River (NCNR). The Aquatic Life Use is impaired for a total of 16.40 miles based on the initial 2008 temperature exceedances and 2012 / 2014 Citizen temperature measurements of the Class V 21°C stockable trout water criterion.

9TOM-1-NCNR (Off Glade Rd. at Heritage Park Trail Lv. 3) Seven temperature measurements exceed the Class V 21°C criterion ranging from at 21.5°C to 26.1°C from 32 measurements within the 2016 and 2014 data windows. Excursions occur during the summer months Lv. 3 [IM]. Two temperature measurements exceed the Class V 21°C criterion at 24.5°C on 7/19/2010 and 24.0°C on 8/19/2010 from 10 measurements for 2012.

9-TOM005.32- (Rt. 725 Bridge upstream of Poverty Creek) One temperature exceedance at 22 C (7/13/17). Temp impairment remains due to upstream and downstream impairment. Both the 2010 and 2008 IRs find two temperature measurements exceed the Class V 21°C criterion from 13 observations. Exceedances occur on 8/15/2005 at 24.4°C and 21.4 °C on 8/15/2006. There are no additional data beyond the 2008 Integrated Report (IR).

9TOM-2-NCNR (Poverty Creek Rd. Bridge Lv. 3) The 2016 data window finds three of 11 temperature measurements exceed the Class V 21°C criterion. Excessive values range from 22 to 24.5°C and occur in the summer months. The 2012 Integrated Report (IR) finds three temperature exceedances of the Class V 21°C criterion occur on 6/16/2010 at 22°C; 7/19/2010 at 24.0°C and 8/16/2010 at 24.5°C from 11 measurements for 2012.

9TOM-3-NCNR (Mt. Zion Road Bridge Lv. 3) Only one temperature measurement (full support) reported during the 2020 data window. Seven temperature measurements exceed the Class V 21°C criterion ranging from at 22.0°C to 24.7°C from 33 measurements within the 2016 data window. Excursions occur during the summer months.

Assessment Unit / Water Name / Location Desc. Cause Category Cause Name VAW-N22R TOM01A00 / Toms Creek / These mainstem waters 5C Temperature		Cycle First Listed 2014	TMDL Dev. Priority	Water Size 4.56
extend from just below the Poverty Creek confluence downstream to the Toms Creek mouth on the New River. These waters are within the WQS five mile public water supply (PWS) designation (NE60).				
VAW-N22R_TOM02A00 / Toms Creek / Toms Creek mainstem 5C Temperature waters just below the Poverty Creek confluence upstream to the mouth of Big Run. These waters are not within the WQS public water supply (PWS) designation (NE60).		2008	L	5.71
VAW-N22R_TOM03A08 / Toms Creek / Toms Creek from the mouth of Big Run upstream to its headwaters (NE60).		2012	L	6.13
Toms Creek	Estuary		servoir	River
Aquatic Life Temperature - Total Impaired Size by Water Type:	(Sq. Miles)	(A	cres)	(Miles) 16.40

Sources:

Source Unknown

New River Basin

Cause Group Code: N22R-05-BAC New River

Cause Location: New River mainstem from Claytor Dam (NE57) downstream to the confluence with Back Creek (NE62).

City / County: Montgomery Co. Pulaski Co. Radford City

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The initial 2016 303(d) Listing of these waters is a result of escherichia coli (E.Coli) excursions of the 235 cfu/100 ml instantaneous criterion in three of 23 samples. All three exceeding samples were found to have E.Coli concentrations of 250 cfu/100 ml. These waters are not meeting the Recreational Use. The 2020 data window extends this Recreational Use impairment upstream to the confluence of Little River.

9-NEW081.72 (Rt. 11 Bridge at Radford) - The 2020 IR finds five of 42 E.coli samples in exceedance of the 235 cfu/100 ml instantaneous criterion.

9-NEW066.90 (New River at Whitethorne) Nine of 36 E.coli samples exceed during the 2020 data window. The 2018 data window shows six of 24 Escherichia coli (E.coli) exceeds the 235 cfu/100 ml instantaneous criterion. Exceedances range from 241 to 383 cfu/100 ml. The 2016 data window found excursions in three of 23 samples. Exceeding values were all 250 cfu/100 ml.

	Cause	9	Cycle First	TMDL Dev.	Water
Assessment Unit / Water Name / Location Desc.	Catego	ry Cause Name	Listed	Priority	Size
VAW-N18R_NEW01A00 / New River / New River mainstem from the Watershed boundary, Crab Creek mouth, upstream to approximately one mile downstream of the Rt. 11 Bridge; end of the WQS public water supply (PWS) section (NE57).		Escherichia coli (E. coli)	2020	L	3.33
VAW-N18R_NEW02A00 / New River / New River mainstem from approximately one mile downstream of the Rt. 11 Bridge upstream the Radford City intake (NE57).		Escherichia coli (E. coli)	2020	L	3.72
VAW-N18R_NEW04A00 / New River / New River mainstem waters from the mouth of Little River upstream to Claytor Dam (NE57).	5A	Escherichia coli (E. coli)	2020	L	0.60
VAW-N22R_NEW02A00 / New River / New River mainstem from the Radford Army Arsenal Plant downstream intake near Whitethorne downstream to the confluence of Back Creek (NE62).	1 5A	Escherichia coli (E. coli)	2016	L	2.86
VAW-N22R_NEW02B14 / New River / New River mainstem from the mouth of Toms Creek downstream to the RAAP downstream intake (NE62).	n 5A	Escherichia coli (E. coli)	2016	L	0.51
VAW-N22R_NEW03A00 / New River / New River mainstem from the confluence of Stroubles Creek downstream to the mouth of Ton Creek (NE59).		Escherichia coli (E. coli)	2016	L	4.09
VAW-N22R_NEW04A00 / New River / New River mainstem from the Radford Army Arsenal Plant upstream intake/Pepper's Ferry Region POTW outfall downstream to the confluence of Stroubles Creek (NE59).	1 5A	Escherichia coli (E. coli)	2016	L	2.32
VAW-N22R_NEW05A00 / New River / New River mainstem from the Blacksburg /Christiansburg /VPI Authority intake at Rt. 114 downstream to the Radford Army Arsenal Plant upstream intake / Pepper's Ferry Regional POTW outfall (NE59).	1 5A	Escherichia coli (E. coli)	2020	L	1.76
VAW-N22R_NEW06A00 / New River / New River mainstem from the Watershed Boundary at the Crab Creek confluence downstrear to the Blacksburg /Christiansburg /VPI Authority intake (NE59).		Escherichia coli (E. coli)	2020	L	1.72

New River Basin

New RiverEstuaryReservoirRiverRecreation(Sq. Miles)(Acres)(Miles)

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

20.91

Sources:

Livestock (Grazing or Feeding Operations) Wildlife Other than

Waterfowl

Unspecified Domestic Waste

Wastes from Pets

Wet Weather Discharges (Non-Point Source)

New River Basin

Cause Group Code: N22R-06-BEN Unnamed Tributaries XEJ and XEH to Slate Branch

Cause Location: Unnamed Tributary XEH from its mouth on Slate Branch upstream to its headwaters. And Unnamed Tributary XEJ

from its mouth on Unnamed Tributary XEH upstream to its headwaters.

City / County: Montgomery Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 5A

The 2008 assessment finds the Aquatic Life Use via the General Standard (Benthic) is impaired for a total of 2.51 miles. Unnamed Tributary to Slate Branch- XEH for 1.68 miles and Unnamed Tributary XEJ to XEH for 0.83 miles. There are no additional data for 9-XEH000.75 and 9-XEJ000.10 beyond the 2008 Integrated Report (IR), 9-XEH000.01 is a new station assessed in 2016.

9-XEH000.75- (Downstream of Villages Development at NRV Mall) There are no additional data beyond the 2008 IR. Bio 'IM' Two 2006 Virginia Stream Condition Index (VSCI) surveys with an average score of 23.1. This station was sampled at the request of the WCRO VWP program with the goal of collecting water quality data prior to new development immediately upstream near the New River Valley Mall complex. A crayfish/macro invertebrate kill in January 2006 impacted the stream with the source occurring somewhere above this station. The most noticeable difference between this site and the reference station is the low abundance of organisms collected in the spring sample compared to the reference site. The abundance increased in the fall and is comparable to the reference site (Falling Branch).

9-XEH000.01 (Near Huckleberry Trail, Downstream of XEJ) - This stream was originally sampled at a location upstream (9-XEH000.75). The 2016 and 2018 data windows find four VSCI scores average 52.0 (2013-2014). The headwaters of Slate Branch are developed with residential and commercial properties as well as Rt. 460 and Peppers Ferry Road. Storm water runoff from these areas may have an impact on water quality at the sampling station which is about one mile downstream of the New River Valley Mall. Habitat scores at this station were relatively good considering the proximity to developed lands upstream and appear favorable for macroinvertebrates. Specific conductance was high at this site during all surveys. Periphyton and algal growth was always thick even during the fall surveys which may be an indication of excessive nutrients.

9-XEJ000.10- (North of NRV Mall) There are no additional data beyond the 2008 IR. Bio 'IM' Two 2006 VSCI surveys with an average score of 23.8. This station was sampled at the request of the WCRO VWP program with the goal of collecting water quality data prior to new development immediately upstream and north of the New River Valley Mall and above the Huckleberry Tail crossing. The main source of impact appears to be recent development and urban land use resulting in altered hydrology. excessive storm water runoff, sediment deposition, bank erosion, and riparian vegetation removal.

Assessment Unit / Water Name / Location Desc.	Caus Catego	e Pry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N22R_XEH01A08 / Slate Branch, UT (XEH) / Unnamed tributary XEH from its mouth on Slate Branch upstream to its headwaters (NE59).	5A	Benthic Macroinvertebra Bioassessments	ates	2008	М	1.68
VAW-N22R_XEJ01A08 / Unnamed Trib. XEJ to XEH / Unnamed Tributary XEJ from its mouth on Unnamed Tributary XEH upstreamits headwaters (NE59).		Benthic Macroinvertebra Bioassessments	ates	2008	M	0.83
Unnamed Tributaries XEJ and XEH to Slate Branch Aquatic Life			Estuary (Sq. Miles)		servoir .cres)	River (Miles)
Benthic Macroinvertebrates Bioassessments - Total	Impaire	d Size by Water Type:				2.51
Sources:						

Loss of Riparian Habitat Municipal (Urbanized High Sediment Resuspension Streambank Density Area) (Clean Sediment) Modifications/Destabilization

New River Basin

Cause Group Code: N23R-01-BAC Sinking Creek

Cause Location: Sinking Creek mainstem waters from just downstream of the Rt. 778 Bridge upstream to the mouth of Gravel Hill

Branch.

City / County: Craig Co. Giles Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The initial 21.03 mile 2010 303(d) Listing of these waters is due to bacteria excursions of the WQS instantaneous criterion for escherichia coli (E.coli).

9-SNK012.06 (Rt. 42 Bridge)- One exceedance of the E.Coli instantaneous criterion (235 cfu/100 ml) occurs within the 2016 and 2018 data windows at 600 cfu/100 ml (2013). The 2010 IR found three of 12 E.coli samples exceed the 235 cfu/100 ml instantaneous criterion. The exceedance range is from 250 to greater than 2000 cfu/100 ml.

9-SNK005.38 (Rt. 778 Bridge)- There are no new data beyond the 2014 data window. E.coli excursions of the 235 cfu/100 ml instantaneous criterion occur in four of 23 observations within the 2014 data window. Values in excess of the instantaneous criterion range from 275 to 600 cfu/100 ml. E.coli excursions of the instantaneous criterion occur in two of 11 observations within the 2010 and 2012 data windows. Values in excess of the instantaneous criterion are 480 and 600 cfu/100 ml.

Assessment Unit / Water Name / Location Desc.	Cause atego	e rry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N23R_SNK01B10 / Sinking Creek / Sinking Creek mainstem waters from just downstream of the Rt. 778 Bridge upstream to the mouth of an unnamed tributary near the Rt. 700 crossing (NE65).	5A	Escherichia coli (E. coli)		2010	Н	3.03
VAW-N23R_SNK01C14 / Sinking Creek / Sinking Creek from just downstream of the Rt. 700 Bridge upstream to the junction of routes 601 & 604 - 6th Order Boundary (NE65)	5A	Escherichia coli (E. coli)		2010	Н	2.74
VAW-N23R_SNK02A00 / Sinking Creek / Sinking Creek from the junction of routes 601 & 604 upstream to the mouth of Gravel Hill Branch- 6th Order Boundary (NE64).	5A	Escherichia coli (E. coli)		2010	Н	15.26
Sinking Creek Recreation			Estuary (Sq. Miles)		servoir cres)	River (Miles)

Sources:

Livestock (Grazing or Feeding Operations)

On-site Treatment Systems

Unspecified Domestic Wet Weather Discharges
(Septic Systems and Similar Waste

Decentralized Systems)

Wet Weather Discharges
(Non-Point Source)

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

21.03

Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N25R-01-BAC Walker Creek

Cause Location: Walker Creek from the Route 52 crossing downstream to the confluence with Kimberling Creek and Town Creek

from the headwaters downstream to the confluence with Crab Orchard Creek.

City / County: Bland Co. Giles Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

Fecal Coliform / 4A

The AWQM station located at 9-WLK060.32 had a 17% and 9-WLK044.06 has 16% exceedance of the E. coli water quality standard and station 9-TNC000.53 had a 55% exceedance of the E. coli water quality standard.

	Cause)		Cycle First	TMDL Dev.	Water
Assessment Unit / Water Name / Location Desc.	Catego	ry Cause Name		Listed	Priority	Size
VAS-N25R_TNC01A10 / Town Creek / From headwaters to Cral Orchard confluence at Town of Bland, WQS Section 1, u (NE66).	b 4A	Escherichia coli (E. coli)	2010	L	4.40
VAS-N25R_WLK04A00 / Walker Creek / Walker Creek mainster from the Kimberling Creek confluence at the Giles/Bland County lir upstream to the Helveys Mill Creek confluence near Point Pleasan WQS Section 1k (NE67).	ne,	Escherichia coli (E. coli)	2006	L	14.48
VAS-N25R_WLK04A12 / Walker Creek / Walker Creek mainster from the Crab Orchard Creek confluence, upstream to the Rt. 52 crossing north of Walker Mountain, WQS Section 1k (NE66).	m 4A	Escherichia coli (E. coli)	2006	L	8.46
VAS-N25R_WLK04B12 / Walker Creek / Walker Creek mainster from the Helveys Mill Creek confluence, near Point Pleasant, upstream to the Crab Orchard Creek confluence, south of Bland, WQS Section 1k (NE67).	m 4A	Escherichia coli (E. coli)	2006	L	10.59
Walker Creek			Estuary	Res	servoir	River
Recreation			(Sq. Miles) (A	cres)	(Miles)
Escherichia coli (E. coli) - Total	Impaired	d Size by Water Type:				37.93
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ory Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N25R_WLK04A00 / Walker Creek / Walker Creek mainster from the Kimberling Creek confluence at the Giles/Bland County lir upstream to the Helveys Mill Creek confluence near Point Pleasan	ne,	Fecal Coliform		2006	L	14.48
WQS Section 1k (NE67).						40.50
	m 4A	Fecal Coliform		2006	L	10.59
WQS Section 1k (NE67). VAS-N25R_WLK04B12 / Walker Creek / Walker Creek mainstel from the Helveys Mill Creek confluence, near Point Pleasant, upstream to the Crab Orchard Creek confluence, south of Bland,	m 4A	Fecal Coliform	Estuary		L	10.59
WQS Section 1k (NE67). VAS-N25R_WLK04B12 / Walker Creek / Walker Creek mainster from the Helveys Mill Creek confluence, near Point Pleasant, upstream to the Crab Orchard Creek confluence, south of Bland, WQS Section 1k (NE67).	m 4A	Fecal Coliform	Estuary (Sq. Miles	Res		

Sources:

Grazing in Riparian or Shoreline Zones

On-site Treatment Systems (Septic Systems and Similar Decentralized Systems) Unrestricted Cattle Access

New River Basin

Cause Group Code: N26R-01-BAC East Wilderness Creek and Nobusiness Creek

Cause Location: This segment includes the mainstem of Nobusiness Creek from the Kimberling Creek confluence upstream 6.4

miles, East Wilderness Creek from the confluence with Wolfpen Branch upstream 3.2 miles.

City / County: Bland Co. Giles Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

AWQM station 9-EWL000.06 had 18% of the samples exceed the E.coli water quality standard. Station 9-NBS000.70 had a

50% exceedance of the E. coli water quality standard.

Escherichia coli (E. coli) -	Total Impaired S	size by Water Type:				10.23
East Wilderness Creek and Nobusiness Creek Recreation			Estuary (Sq. Miles)		servoir cres)	River (Miles)
VAS-N26R_NBS01B04 / Nobusiness Creek / Nobusiness Grom Kimberling Creek confluence to upstream of Panther De Branch in WQS Section 1, DGIF *** (NE69).		scherichia coli (E. coli)		2010	L	6.72
VAS-N26R_EWL01A10 / East Wilderness Creek / A Wolfp Branch tributary near Shady Grove Church from Wolf Creek Mountain to the north, WQS Section 1 (NE68).	en 4A E	scherichia coli (E. coli)		2010	L	3.51
Assessment Unit / Water Name / Location Desc.	Cause Category	Cause Name		Cycle First Listed	Dev. Priority	Water Size

Sources:

Grazing in Riparian or Shoreline Zones

Unrestricted Cattle Access

New River Basin

Cause Group Code: N26R-03-TEMP Nobusiness Creek

Cause Location: This segment includes from the Kimberling Creek confluence to upstream of Panther Den Branch.

City / County: Bland Co. Giles Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5A

AWQM station 9-NBS000.70 had a 18% exceedance of the WQS for temperature in Class VI waters.

Cause Cycle TMDL
First Dev. Water
Assessment Unit / Water Name / Location Desc. Category Cause Name Listed Priority Size

VAS-N26R_NBS01B04 / Nobusiness Creek / Nobusiness Creek 5A Temperature 2018 L 6.72

from Kimberling Creek confluence to upstream of Panther Den

Branch in WQS Section 1, DGIF *** (NE69).

Nobusiness Creek
Aquatic Life
Estuary Reservoir River
(Sq. Miles) (Acres) (Miles)

Temperature - Total Impaired Size by Water Type: 6.72

Sources:

Source Unknown

New River Basin

Cause Group Code: N27R-01-BAC Little Walker Creek

Cause Location: Little Walker Creek mainstem from its confluence with Walker Creek upstream to the mouth of Spur Branch.

City / County: Pulaski Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The initial 2004 303(d) Listing of these waters is the result of fecal coliform (FC) bacteria exceedances (two exceeding from 18 observations) causing a 17.48 mile impairment. Escherichia coli (E.coli) replaces fecal coliform bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters].

9-LWK000.77 (Rt. 100 Bridge) Escherichia coli (E.coli) exceed the 235 cfu/100 ml instantaneous criterion in eight of 34 and five of 22 samples within the 2018 and 2016 data windows, respectively. Exceedance range is 337 to greater than 2000 cfu/100 ml. The 2014 data window found three of 11 samples in exceedance of the instantaneous criterion. Values exceeding the criterion range from 275 to greater than 2000 cfu/100 ml. The 2008 through 2012 assessments find E.coli exceed the instantaneous criterion in five of 12 samples. Values exceeding the criterion range from 320 to 500 cfu/100 ml. Four of nine excursions are reported in 2006 with the range of exceedance from 350 to 500 cfu/100 ml.

Cycle TMDI

Assessment Unit / N	Water Name /	Location Desc.	Cause Catego	e ry Cause Name	First Listed	Dev. Priority	Water Size
VAW-N27R_LWK01A00 mainstem from its conflu				Escherichia coli (E. coli)	2006	Н	17.48

mouth of Spur Branch (NE72).

Little Walker Creek	Estuary	Reservoir	River
Recreation	(Sq. Miles)	(Acres)	(Miles)
Esche	erichia coli (E. coli) - Total Impaired Size by Water Type:		17.48

Sources:

Livestock (Grazing or **On-site Treatment Systems Unspecified Domestic** Wet Weather Discharges (Non-Point Source) Feeding Operations) (Septic Systems and Similar Waste Decentralized Systems)

Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N29R-01-PCB New River, Reed Creek, Claytor Lake, Peak Creek, Stony Creek, Walker Creek, And Unknown Tributary XAG.

Cause Location: The impairment begins at the I-77 bridge crossing the New River and extends downstream to the VA/WVA State Line and includes the tributaries Peak Creek and Reed Creek as described below.

City / County: Giles Co. Montgomery Co. Pulaski Co. Radford City Wythe Co.

Use(s): Fish Consumption Wildlife

Cause(s) / VA Category: PCBs in Fish Tissue / 4A Polychlorinated Biphenyls (PCBs) / 4A

The VDH issued a fish consumption advisory 8/6/2001 for PCBs for the lower portion of the New River (Rt. 114 Bridge downstream to the VA/WV Line - 52.0 mi) based on fish tissue collections from carp. Advisory extension to Claytor dam was issued 8/6/2003 (11.47 mi) and further extended upstream on New River (13 mi) to the I-77 Bridge to include the lower portions of Peak Ck (4.02 mi), Reed Ck (16.35 mi) and Claytor Lake (4,287 ac) on 12/2/2004. Stony Ck is a 2010 IR addition to the original 2002 303(d) Listing. An unnamed tributary (XAG) is an addition with the 2016 IR. The VDH level of concern is 50 parts per billion (ppb) in fish tissue.

Water column (WC)data from 2010-2016 are listed below where excursions of the WQS criterion of 640 pg/L are contravened causing an Observed Effect (OE) or 303(d) Listing for 'PCBs in Water Column'. Collections are made in wet weather (WW) & dry weather (DW) conditions.

A complete listing of fish tissue sites and data are available at https://www.deq.virginia.gov. VDH Advisory information is available at https://www.vdh.virginia.gov.

9-RDC009.00 (Near Rt. 619 at Grahams Forge) 2012 2 species: Carp 5 fish composite [62.6-69.4 cm] at 68.24 ppb. Smouth Bass 5 fish comp. [21.8-26.6] 3.04 ppb. 2014 2 sp: Carp 4 fish [67.9-76.5] 75.67 ppb & 5 fish [64.5-69.8] 85.77. Smouth Bass 5 fish [23.1-30.3] 2.46 ppb.

9-NEW107.51 (New R. near Allisonia) 2014 3 sp: Channel Catfish 3 fish comp. [61.2-69.5] 23.02 ppb; Smouth Bass 3 fish [40.1-49.6] 2.45 ppb; Carp 5 fish [56.5-70.4] 45.12 ppb & 5 fish [55.3-71.3] 8.79 ppb.

9-NEW098.32 (Rt. 672 Br, Lighthouse) 2012 4 sp: Channel Catfish 2 fish [70.5-71.5] 65.15 ppb. Lmouth Bass 5 fish [34.5-43.1] 7.76 ppb; Spotted Bass 5 fish [34.2-38.2] 11.00 ppb; Carp 3 fish [45.8-56.5] 6.04 ppb.

9-PKC009.53 (Upstream of XAG confluence) 2014 DW 197.02 pg/L- 'FS'; WW 799.75 pg/L- 'OE'

9-XAG000.01 (Mouth of X-Trib XAG near former Allied Site) 2014: DW 1,458.87 pg/L (7/31/2014); WW 1,754.02 pg/L (10/15/2014).

9-XFQ000.77 (Off Pierce Ave. near Calfee Park) 2014: DW 194.99 pg/L- 'FS'; WW 686.76 pg/L- 'OE'

9-PKC007.82 (Rt. 99 Br) 2012 3 sp: Stoneroller 15 fish [14.3-16.0] 33.18 ppb. Rock Bass 5 fish [16.7-18.6] 10.49 ppb; Redbreast Sunfish 5 fish [14.3-18.1] 3.01 ppb.

2013: DW 1,193.64 pg/L; WW 2,436.73 pg/L; 2014 1 sample 'FS DW 389.51 pg/L; WW 1,252.42 pg/L

9-PKC004.65 (Rt. 100 Br) 2012 4 sp: Channel catfish2 fish [63.1-69.0] 33.15 ppb. Lmouth Bass 5 fish [33.4-40.8] 2.68 ppb; Carp 2 sizes: 4 fish [54.6-62.0] 2.32 ppb & 4 fish [54.6-62.0] 9.16 ppb; Smouth Bass 3 fish [35.3-42.6] 6.90 ppb. 2014 WC PCB: 1,075.73 pg/L

9-NEW088.86 (New R. Claytor Lake at Dam) 2012 6 sp: Flathead Catfish 2 fish [83.0-87.5] 86.67 ppb. Carp 4 fish [56.5-67.0] 2.05 ppb; Channel Catfish 1 fish [58.8 cm] 7.43 ppb; Lmouth Bass 5 fish [32.5-34.5] 0.36 ppb; Smouth Bass 4 fish [27.0-32.2] 0.88 ppb; Spotted Bass 3 fish [28.8-36.8] 0.00 ppb.

9-NEW085.94 (New R. downstream of Claytor Dam) 2012 2 sp: Flathead Catfish 5 fish [57.5-70.3] 33.74 ppb. Carp 5 fish [62.6-81.0] 11.27 ppb.

9-NEW081.72 (Rt. 11 Br at Radford) 2010: DW 320 pg/L - 'FS'; WW 4,739- 'OE'. 2011 WW 243.70 pg/L - 'FS'. 2013 WW 647.88 pg/L- 'OE'

New River Basin

9-NEW079.19 (New R. below Radford Univ.) 2012 1 sp: Carp 2 fish [67.5-76.5] 53.28 ppb & 2 fish [76.8-83.6] 94.85 ppb.

9-NEW066.90 (New R. at Whitethorne) 2012 1 sp: Carp 1 fish [72.0 cm] 125.58 ppb.

9-LWK000.77 (Rt. 100 Br) 2011 WW 642.4 pg/L- 'OE'; 2014 DW 39.79 pg/L- 'FS'

9-WLK004.34 (Rt. 622 Br - Giles Co.) 2010 WW 1,706 pg/L & 2011 WW 648.8 pg/L. 2014 DW 60.12 pg/L 'FS'

9-NEW050.70 (New R. near Pembroke) 2012 3 sp: Carp 2 fish [67.5-71.6] 419.87 ppb; Channel Catfish 1 fish [58.1] 23.26 ppb. Flathead Catfish 1 fish [51.4] 9.60 ppb.

9-SNC000,20- 2004 3 sp; Rock Bass 10 fish [16-20] 25.21, Smouth Bass 3 fish [28,6-30,5] 22.13, White sucker 1 fish [38,4] 0.08 ppb.

9-NEW038.71 (New R. below Celeanse) 2012 2 sp: Carp 2 fish [68.1-69.0] 355.63 ppb; Flathead Catfish 1 fish [56.0] 25.39 ppb. 2010 DW- 129 pg/L- 'FS; WW 784 pg/L- 'OE' and 2011 WW- 222 pg/L- 'FS'

9-WFC003.69 (Rt. 724 Br at Gage) 2014 DW 114.7 pg/L- 'FS'; WW 1,527.45 pg/L- 'OE'

9-WFC000.20 (Rt. 61 Br) 2011 WW 1,220.1 pg/L- 'OE'; 2013 WW 201.31 pg/L- 'FS'. 2014 WW 163.39 pg/L- 'FS'; DW 117.93 pg/L- 'FS'.

9-NEW030.15 (Rt. 460 Br at Glen Lyn) 2012 1 sp: Carp 1 fish [85.0] 234.01 ppb; Carp 2 fish [72.5-74.8] 448.15 ppb.

9-NEW031.00 (Abv Glen Lyn) 2010 DW- 66 pg/L 'FS'; WW- 841 pg/L 'OE'

9-NEW028.95 (New R. below Glen Lyn) 2010 DW 177.5 pg/L- 'FS'; WW- 709.9 pg/L 'OE'. 2011 DW- 110.4 pg/L 'FS'; WW-399.6 pg/l 'FS'. 2013 WW 284.08 pg/L- 'FS'; NA 116.76 pg/L- 'FS'. 2014 WW 529.48 pg/L- 'FS'; DW 121.59 pg/L- 'FS'

Assessment Unit / Water Name / Location Desc.	Cause Catego	e rry Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size	
VAS-N08R_NEW01A02 / New River / Mainstem, north of Barren Springs, from Reed Creek confluence downstream to Big Reed Island Creek confluence, WQS Section 2 (NE43).	4A	PCBs in Fish Tissue	2006	L,	5.71	
VAS-N11R_RDC01B00 / Reed Creek / Lower mainstem from Muskrat Branch confluence downstream to Rt. 52 bridge south of Max Meadows, WQS Section 2 (NE29).	4A	PCBs in Fish Tissue	2016	L	5.85	
VAS-N11R_RDC01B06 / Reed Creek / Lower mainstem from Rt. 52 bridge downstream to Miller Creek confluence south of Max Meadows, WQS Section 2 (NE31).	4A	PCBs in Fish Tissue	2006	L	0.60	
VAS-N11R_RDC02B02 / Reed Creek / Reed Creek from Miller Creek confluence at Max Meadows downstream to the Glade Creek confluence, near Boiling Spring, WQS Section 2g (NE31).	4A	PCBs in Fish Tissue	2006	L	6.08	
VAS-N11R_RDC03B04 / Reed Creek / From New River confluence near Lone Ash, upstream to the Glade Creek confluence near Boiling Spring, WQS Section 2 (NE31).	4A	PCBs in Fish Tissue	2006	L	9.87	
VAW-N16L_NEW01A02 / Claytor Lake (New River) / Claytor Lake from its impounding structure upstream to the Claytor State Park Cabins.	4A	PCBs in Fish Tissue	2006	L 1	,196.91	
VAW-N16L_NEW01B14 / Claytor Lake (New River) / Claytor Lake from the Claytor State Park Cabins upstream to the former Burlingtor Industries water intake.		PCBs in Fish Tissue	2006	L	602.03	
VAW-N16L_NEW02A02 / Claytor Lake (New River) / Claytor Lake from the Claytor State Park Cabins upstream to the confluence of	4A	PCBs in Fish Tissue	2006	L	278.51	
Final 2020 Appendi	ix 5 - 34	189				

New River Basin					
Peak Creek					
VAW-N16L_NEW03A02 / Claytor Lake (New River) / Claytor Lake from the confluence of Peak Creek upstream to the end of the WQS public water supply (PWS) designation. The segment ends five miles upstream of the former Burlington Industries intake.	4A	PCBs in Fish Tissue	2006	L	671.89
VAW-N16L_NEW04A02 / Claytor Lake (New River) / Claytor Lake from the end of the Burlington WQS public water supply (PWS) designation upstream to the Pulaski County PSA intake.	4A	PCBs in Fish Tissue	2006	L	447.80
VAW-N16L_NEW05A02 / Claytor Lake (New River) / Claytor Lake from the Pulaski County PSA intake upstream to the end of the WQS public water supply (PWS) designation. Five miles upstream from the Pulaski County PSA intake.	4A	PCBs in Fish Tissue	2006	L	660.27
VAW-N16L_NEW06A02 / Claytor Lake (New River) / Claytor Lake from the upstream end of the Pulaski County PSA WQS public water supply (PWS) designation upstream to the backwaters of Claytor Lake at Allisonia.	4A	PCBs in Fish Tissue	2006	L	152.13
VAW-N16R_NEW01A00 / New River / This section of the New River extends from the mouth of Big Reed Island Creek downstream to the backwaters of Claytor Lake Class IV sec. 2c (NE43).	4A	PCBs in Fish Tissue	2006	L	0.61
VAW-N17L_PKC01A10 / Claytor Lake (Peak Creek) / Peak Creek from its confluence with the New River upstream to the end of the WQS public water supply (PWS) designation.	4A	PCBs in Fish Tissue	2002	L	216.86
VAW-N17L_PKC02A10 / Claytor Lake (Peak Creek) / Peak Creek from the end of the WQS public water supply (PWS) designation upstream to its backwaters.	4A	PCBs in Fish Tissue	2002	L	78.16
VAW-N17R_PKC01A00 / Peak Creek / This portion of Peak Creek begins just downstream of the Rt. 99/Norfolk Southern crossing extending downstream to the inundation of Peak Creek in Claytor Lake (NE46).	4A	PCBs in Fish Tissue	2002	L	1.83
VAW-N17R_PKC02A00 / Peak Creek / The segment begins downstream of the Washington Ave. Bridge (~0.20 miles) and extends on downstream to just below the Rt. 99 Bridge/Norfolk Southern Railway crossing of Peak Creek (NE46).	4A	PCBs in Fish Tissue	2002	L	1.66
VAW-N17R_PKC03A00 / Peak Creek / This portion of Peak Creek extends from the mouth of Tract Fork to downstream of the Washington Ave. Bridge (~0.20 miles) (NE46).	4A	PCBs in Fish Tissue	2006	L	0.51
VAW-N18R_NEW01A00 / New River / New River mainstem from the Watershed boundary, Crab Creek mouth, upstream to approximately one mile downstream of the Rt. 11 Bridge; end of the WQS public water supply (PWS) section (NE57).	4A	PCBs in Fish Tissue	2006	L	3.33
VAW-N18R_NEW02A00 / New River / New River mainstem from approximately one mile downstream of the Rt. 11 Bridge upstream to the Radford City intake (NE57).	4A	PCBs in Fish Tissue	2006	L	3.72
VAW-N18R_NEW03A00 / New River / New River mainstem from the City of Radford water intake upstream to the confluence of Little River (NE57).	4A	PCBs in Fish Tissue	2006	L	2.14
VAW-N18R_NEW04A00 / New River / New River mainstem waters from the mouth of Little River upstream to Claytor Dam (NE57).	4A	PCBs in Fish Tissue	2006	L	0.60
VAW-N22R_NEW01A00 / New River / The New River mainstem from the confluence of Back Creek downstream to the Watershed Boundary at the Montgomery / Giles County Line (NE62).	4A	PCBs in Fish Tissue	2002	L	3.44

New River Basin					
VAW-N22R_NEW02A00 / New River / New River mainstem from the Radford Army Arsenal Plant downstream intake near Whitethorne downstream to the confluence of Back Creek (NE62).	4A	PCBs in Fish Tissue	2002	L	2.86
VAW-N22R_NEW02B14 / New River / New River mainstem from the mouth of Toms Creek downstream to the RAAP downstream intake (NE62).	4A	PCBs in Fish Tissue	2002	L	0.51
VAW-N22R_NEW03A00 / New River / New River mainstem from the confluence of Stroubles Creek downstream to the mouth of Toms Creek (NE59).	4A	PCBs in Fish Tissue	2002	L	4.09
VAW-N22R_NEW04A00 / New River / New River mainstem from the Radford Army Arsenal Plant upstream intake/Pepper's Ferry Region POTW outfall downstream to the confluence of Stroubles Creek (NE59).	4A	PCBs in Fish Tissue	2002	L	2.32
VAW-N22R_NEW05A00 / New River / New River mainstem from the Blacksburg /Christiansburg /VPI Authority intake at Rt. 114 downstream to the Radford Army Arsenal Plant upstream intake / Pepper's Ferry Regional POTW outfall (NE59).	4A	PCBs in Fish Tissue	2002	L	1.76
VAW-N22R_NEW06A00 / New River / New River mainstem from the Watershed Boundary at the Crab Creek confluence downstream to the Blacksburg /Christiansburg /VPI Authority intake (NE59).	4A	PCBs in Fish Tissue	2006	L	1.72
VAW-N23R_NEW01A00 / New River / New River mainstem from the Giles/Montgomery County Line downstream to the confluence of Sinking Creek (NE63).	4A	PCBs in Fish Tissue	2002	L	5.47
VAW-N24R_NEW01A00 / New River / New River mainstem from the confluence of Stony Creek upstream to the mouth of Walker Creek on the New River (NE74).	4A	PCBs in Fish Tissue	2002	L	3.87
VAW-N24R_NEW02A00 / New River / New River mainstem waters from the mouth of Walker Creek upstream to the confluence of Little Stony Creek with the New River (NE74).	4A	PCBs in Fish Tissue	2002	L	2.00
VAW-N24R_NEW03A00 / New River / New River mainstem waters from the confluence of Little Stony Creek upstream to mouth of Sinking Creek on the New River. (NE74)	4A	PCBs in Fish Tissue	2002	L	3.87
VAW-N28R_SNC01A00 / Stony Creek / Stony Creek mainstem waters from its mouth on the New River upstream to Chemical Lime Company's outfall on Stony Creek (NE75).	4A	PCBs in Fish Tissue	2010	L	1.36
VAW-N28R_SNC02A00 / Stony Creek / Stony Creek mainstem waters from the Chemical Lime Company outfall on Stony Creek upstream to the Kimballton Branch confluence on Stony Creek (NE75).	4A	PCBs in Fish Tissue	2010	L	0.63
VAW-N28R_SNC03A00 / Stony Creek / Stony Creek mainstem waters from the confluence of Kimballton Branch upstream to the mouth of Laurel Branch (NE75).	4A	PCBs in Fish Tissue	2010	L	1.69
VAW-N28R_SNC04A00 / Stony Creek / Stony Creek mainstem from the confluence of Laurel Branch upstream to the mouth of Pine Swamp Branch (NE75).	4A	PCBs in Fish Tissue	2010	L	4.69
VAW-N29R_NEW01A02 / New River / New River mainstem from the backwaters of Bluestone Reservoir, Route 460, to the confluence of Rich Creek.	4A	PCBs in Fish Tissue	2002	L	3.20
VAW-N29R_NEW02A02 / New River / New River mainstem from the mouth of Rich Creek upstream to the confluence of Wolf Creek.	4A	PCBs in Fish Tissue	2002	L	3.55
VAW-N29R_NEW03A02 / New River / New River mainstem from the confluence of Wolf Creek upstream to the Celanese Acetate	4A	PCBs in Fish Tissue	2002	L	2.79
Final 2020 Appendix	5 - 34	191			

New River Basin						
Plant outfalls.						
VAW-N29R_NEW04A02 / New River / New River mainstem from the Celeanse Acetate Plant outfalls upstream to the watershed boundary at the confluence of Stony Creek.	4A	PCBs in Fish Tissue		2002	L	5.78
VAW-N35R_NEW01A00 / New River / New River mainstem from the Rt. 460 Bridge at Glen Lyn downstream to the Virginia/West Virginia State Line.	4A	PCBs in Fish Tissue		2002	L	6.92
New River, Reed Creek, Claytor Lake, Peak Creek, Stony Creek, Wa Unknown Tributary XAG. Fish Consumption	alker	Creek, And	Estuary (Sq. Miles)		servoir cres)	River (Miles)
PCBs in Fish Tissue - Total Imp	paired	d Size by Water Type:		4,3	304.56	105.03
•	Cause atego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAW-N17R_PKC01A00 / Peak Creek / This portion of Peak Creek begins just downstream of the Rt. 99/Norfolk Southern crossing extending downstream to the inundation of Peak Creek in Claytor Lake (NE46).	4A	Polychlorinated Bipheny	ls (PCBs)	2016	L	1.83
VAW-N17R_PKC02A00 / Peak Creek / The segment begins downstream of the Washington Ave. Bridge (~0.20 miles) and extends on downstream to just below the Rt. 99 Bridge/Norfolk Southern Railway crossing of Peak Creek (NE46).	4A	Polychlorinated Bipheny	ls (PCBs)	2016	L	1.66
VAW-N17R_XAG01A02 / Peak Creek, UT (XAG) / An unnamed tributary to Peak Creek not within WQS designated public water supply (PWS) sections. The unnamed tributary mouth is located @37°02'47" / 80°46'03" (NE46).	4A	Polychlorinated Bipheny	ls (PCBs)	2016	L	3.20
VAW-N25R_WLK01A00 / Walker Creek / Walker Creek mainstem waters from its mouth on the New River upstream to the Cecil Branch confluence at the Rt. 100 crossing (NE73).	4A	Polychlorinated Bipheny	ls (PCBs)	2014	L	8.39
New River, Reed Creek, Claytor Lake, Peak Creek, Stony Creek, Wa Unknown Tributary XAG. Fish Consumption	alker	Creek, And	Estuary (Sq. Miles)		servoir .cres)	River (Miles)
Polychlorinated Biphenyls (PCBs) - Total Imp	oaired	d Size by Water Type:				15.08

Discharges from Municipal Industrial Point Source Separate Storm Sewer Systems (MS4)

Source Unknown

Industrial Point Source Stormwater Discharge (Permitted)

Non-Point Source Stormwater Discharge (Permitted)

New River Basin

Cause Group Code: N30R-01-BAC Wolf Creek and Tributaries

Cause Location: This segment extends from the Burkes Garden Creek confluence downstream between the confluence with Clear

Fork and Wilderness Creek and Little Creek, a Wolf Creek tributary upstream to the Tazewell County Sportsmen

Club impoundment.

City / County: Bland Co. Tazewell Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A Fecal Coliform / 4A

The AWQM station located at 9-WFC039.16 had a 91% exceedance of the E.coli water quality standard. Station 9-WFC050.16 had a 78% exceedance, 9-WFC032.47 had 41%, and station 9-WFC024.57 had a 17% exceedance. Station 9-LTL001.22 had a 50% exceedance of the E. coli water quality standard.

	Cause			Cycle First	TMDL Dev.	Water
Assessment Unit / Water Name / Location Desc. Ca	ategor	y Cause Name		Listed	Priority	Size
VAS-N30R_LTL02A10 / Little Creek / A Wolf Creek tributary upstream to Tazewell County Sportsmen Club impoundment in WQS Section 1 (NE78).	4A	Escherichia coli (E. coli)		2014	L	1.89
VAS-N30R_WFC01A00 / Wolf Creek / Mainstem from unnamed tributary downstream of Carter Branch at Grapefield downstream to the Hunting Camp Creek confluence north of Bastian, WQS Section 1 (NE78).	4A	Escherichia coli (E. coli)		2006	L	9.11
VAS-N30R_WFC01A04 / Wolf Creek / From Burkes Garden Creek confluence downstream to unnamed tributary downstream of Carter Branch at Grapefield, WQS Section 1, DGIF *** (NE78).	4A	Escherichia coli (E. coli)		2006	L	7.97
VAS-N30R_WFC01A06 / Wolf Creek, headwaters / Upper segment of Wolf Creek inside Burkes Garden from Snyder Branch confluence downstream to Little Creek confluence (37.1484/-81.2483), WQS Section 1 (NE76/78)	4A	Escherichia coli (E. coli)		2006	L	3.80
VAS-N30R_WFC01B06 / Wolf Creek / Mainstem from the Hunting Camp Creek confluence downstream to Wilderness Creek confluence at South Gap, WQS Section 1 (NE78).	4A	Escherichia coli (E. coli)		2006	L	6.39
VAS-N32R_WFC01A10 / Wolf Creek / Wolf Creek between confluence with Clear Fork at Rocky Gap and Wilderness Creek at South Gap, WQS Section 1, u, parallel to I-77 at Rocky Gap (NE78).	4A	Escherichia coli (E. coli)		2010	L	1.89
Wolf Creek and Tributaries			Estuary	- D	servoir	River
Recreation			(Sq. Miles)		cres)	(Miles)
Recreation Escherichia coli (E. coli) - Total Imp	paired	Size by Water Type:				(Miles) 31.05
Escherichia coli (E. coli) - Total Imp	Cause	Size by Water Type: y Cause Name				,
Escherichia coli (E. coli) - Total Imp	Cause ategor			Cycle First	TMDL Dev.	31.05 Water
Escherichia coli (E. coli) - Total Imp Assessment Unit / Water Name / Location Desc. VAS-N30R_WFC01A00 / Wolf Creek / Mainstem from unnamed tributary downstream of Carter Branch at Grapefield downstream to the Hunting Camp Creek confluence north of Bastian, WQS Section	Cause ategor	y Cause Name		Cycle First Listed 2002	TMDL Dev. Priority	31.05 Water Size

New River Basin

Sources:

Grazing in Riparian or Shoreline Zones

Unrestricted Cattle Access

New River Basin

Cause Group Code: N30R-01-BEN Little Creek

Cause Location: This segment includes the mainstem from the confluence with Wolf Creek upstream to the Tazewell County

Sportsmen's' Club impoundment.

City / County: Bland Co. Tazewell Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 5A

Benthic station 9-LTL001.22 was impaired based on VSCI scores of 58 in 2011 and 52 and 64 in 2012.

TMDL Cycle First Dev. Cause Water Listed **Priority** Size Category Cause Name Assessment Unit / Water Name / Location Desc. VAS-N30R_LTL02A10 / Little Creek / A Wolf Creek tributary 5A Benthic Macroinvertebrates 2010 Н 1.89 Bioassessments

upstream to Tazewell County Sportsmen Club impoundment in WQS Section 1 (NE78).

Little Creek

Aquatic Life

Estuary Reservoir River
(Sq. Miles) (Acres) (Miles)

1.89

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type:

Sources:

Grazing in Riparian or Loss of Riparian Habitat Unrestricted Cattle Access Shoreline Zones

New River Basin

Cause Group Code: N31R-01-BAC Hunting Camp Creek

Cause Location: This segment extends from the confluence with Wolf Creek, upstream through the community of Bastian to an

impoundment, river mile 8.50.

City / County: Bland Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The AWQM station located at 9-HCC001.40 had a 18% exceedance and station 9-HCC005.57 had a 58% exceedance of the

bacteria water quality standard.

Cause Cycle TMDL
Cause First Dev. Water
Assessment Unit / Water Name / Location Desc. Category Cause Name Listed Priority Size
AS-N31R HCC01A00 / Hunting Camp Creek / Segment is from 4A Escherichia coli (E. coli) 2006 L 8.93

VAS-N31R_HCC01A00 / Hunting Camp Creek / Segment is from the confluence with Wolf Creek, upstream through the community of Bastian to an impoundment, river mile 8.50, WQS Section 1, DGIF *** (NE77).

Hunting Camp Creek

Recreation

Estuary Reservoir River
(Sq. Miles) (Acres) (Miles)

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

8.93

Sources:

Crop Production (Crop Land Livestock (Grazing or

or Dry Land) Feeding Operations)

Loss of Riparian Habitat

New River Basin

Cause Group Code: N32R-01-BAC Wolf Creek

Cause Location: Wolf Creek mainstem waters from the mouth of Clear Fork Creek downstream to the confluence of Wolf Creek with

the New River.

City / County: Bland Co. Giles Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The originally listed 2004 portion of the overall extent described above began near the intersection of Routes 61 and 724 at the confluence of an unnamed tributary extending downstream to the mouth of Wolf Creek on the New River. A total of 5.60 miles. A bacteria TMDL was completed in 2015: E.coli TMDL Development for Wolf Creek and Tributaries in Giles, Bland, and Tazewell Counties, VA [Approved: EPA 7/27/16, SWCB 6/27/16; TMDL ID: 66175].

The 2006 Integrated Report (IR) extends the 2004 303(d) Listed fecal coliform (FC) bacteria impairment 16.71 miles upstream. The total bacteria impairment is 22.31 miles. Escherichia coli (E.coli) replaces fecal coliform (FC) bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters].

9-WFC017.31 (Bridge #6065 on Rt. 644 off Rt. 61) Six out of 36 E.Coli samples exceed the 235 cfu/100 ml instantaneous criterion within the 2016 data window. Exceedances range from 250 to greater than 2000 cfu/100 ml. Five exceeding values are found from 24 escherichia coli (E.coli) observations in 2014. The range of exceedance is from 250 to greater than 2000 cfu/100 ml. Two of 15 E.coli samples exceed the 235 cfu/100 ml criterion at 300 and 450 cfu/100 ml within the 2012 data window. The 2010 and 2008 assessments find two of 12 E.coli samples exceed the 235 cfu/100 ml criterion at 420 and 1000 cfu/100 ml. Two of nine E.coli samples exceed the criterion with the same exceedances in 2006.

9-WFC011.05- (Rt. 676 Bridge at Boxely) The 2016 data window finds two of 36 E.Coli samples exceed the 235 cfu/100 ml instantaneous criterion (400 cfu/100ml and >2000 cfu/100 ml). One exceeding value of greater than 2000 cfu/100 ml exceeds the 235 cfu/100 ml instantaneous criterion from 12 observations within the 2014 data window. There were no additional data within the 2012 data window. Both the 2008 and 2010 assessments find E.coli bacteria exceed the instantaneous criterion in two of 10 samples. Exceeding values are both at 700 cfu/100 ml.

9-WFC005.61 (Rt. 673 Bridge at Penvir) Four of 35 E.Coli samples exceed the instantaneous criterion of 235 cfu/100 ml in the 2016 data window. Excursions range from 250 to 625 cfu/100 ml. The 2014 data window found two of 12 samples in exceedance of the 235 cfu/100 ml instantaneous criterion. Excessive values are 300 and 625 cfu/100 ml. There were no additional data within the 2012 data window. E.coli exceedances are found in five of 12 samples in 2008 and 2010. Values in excess of the 235 cfu/100 ml criterion range from 250 to greater than 2000. E.coli exceedances are found in three of nine samples and the same range of exceedance as in 2008.

9-WFC000.20 (Rt. 61 Bridge) E.coli samples exceed the 235 cfu/100 ml instantaneous criterion in five of 48 samples within the 2018 data window. The 2016 data window finds three of 48 samples exceed the 235 cfu/100 ml instantaneous criterion. Excursions range from 275 to 425 cfu/100 ml. Two of 24 escherichia coli (E.coli) samples exceed the 235 cfu/100 ml instantaneous criterion in 2014. The 2012 assessment finds one of 14 E.coli samples exceeding the instantaneous criterion of 235 cfu/100 ml at 1200. E.coli exceeds the instantaneous criterion in three of 12 samples in 2008 and 2010. Each excursion of the criterion is 520, 900 and 1200 cfu/100 ml. E.coli excursions in 2006 are two of nine samples with each excursion of the criterion at 520 and 900 cfu/100 ml. The 2004 Integrated Report (IR) finds FC exceedances of the 400 cfu/100 ml instantaneous criterion in two of 18 samples resulting in a 2004 impairment listing that remains. Exceeding values are 700 and 1500 cfu/100 ml.

Cycle TMDI

_	Causo	e ory Cause Name	First Listed	Dev. Priority	Water Size	
VAW-N32R_WFC01A00 / Wolf Creek / Wolf Creek mainstem from its mouth on the New River upstream to the former Narrows STP outfall on Wolf Creek. Mill Creek confluence (NE81)	4A	Escherichia coli (E. coli)	2006	L	0.39	
VAW-N32R_WFC02A00 / Wolf Creek / Wolf Creek mainstem from the mouth of Mill Creek former Narrows STP outfall upstream to an unnamed bridge crossing Wolf Creek (NE81).	4A	Escherichia coli (E. coli)	2006	L	5.22	
VAW-N32R_WFC03A00 / Wolf Creek / Wolf Creek mainstem	4A	Escherichia coli (E. coli)	2006	L	8.80	
Final 2020 Appendix	c 5 - 3	497				

New River Basin

waters from an unnamed bridge upstream to Bland/Giles County Line (NE81).

VAW-N32R_WFC04A00 / Wolf Creek / Wolf Creek mainstem waters from the Bland/Giles County Line upstream to the confluence of Clear Fork Creek (NE81).

4A Escherichia coli (E. coli)

2006

7.91

Wolf Creek
Recreation

Estuary (Sq. Miles)

Reservoir (Acres)

River (Miles)

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

22.32

Sources:

Livestock (Grazing or Feeding Operations)

Municipal (Urbanized High

Density Area)

On-site Treatment Systems (Septic Systems and Similar Decentralized Systems)

Unspecified Domestic Waste

Wastes from Pets

Wet Weather Discharges (Non-Point Source)

Wildlife Other than Waterfowl

New River Basin

Cause Group Code: N32R-01-TEMP Wolf Creek

Cause Location: Wolf Creek mainstem waters from the Bland/Giles County Line upstream to the confluence of Clear Fork Creek.

City / County: Bland Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5C

The Aquatic Life Use impairment for temperature returns with the 2014 Integrated Report (IR).

9-WFC017.31 (Bridge #6065 on Rt. 644 off Rt. 61) No additional data collected since the 2016 data window. The 2016 data window finds four of 36 exceedances of the 21°C Class V - Stockable Trout Waters criterion. One exceedance occurs at 22.9°C (7/23/13) in addition to the exceedances within the 2014 data window. Three of 24 temperature measurements exceed the WQS Class V - Stockable Trout water criterion of 21°C in 2014. Values in excess of the criterion are 24.7°C (7/12/2011), 23.0°C (6/25/2012) and 21.2°C (8/21/2012). These waters were delisted with the 2012 IR as temperature excursions of the WQS Class V criterion of 21°C are zero of 15 measurements or an exceedance rate of 0.0% at station 9-WFC017.31. Originally listed in 2008 these waters should have been listed in 2006 with two of nine exceeding values and a TMDL Schedule of 2018. Two of 12 temperature measurements exceed the Class V stockable trout water 21°C criterion within the 2008 and 2010 data windows. Exceeding values are 21.1°C on 8/4/2003 and 21.9°C on 8/30/2004.

Assessment Unit / Water Name / Location Desc.	Cause Category Cause Name	First Listed	Dev. Priority	Water Size
VAW-N32R_WFC04A00 / Wolf Creek / Wolf Creek mainstem waters from the Bland/Giles County Line upstream to the confluence.	5C Temperature	2014	М	7.91
of Clear Fork Creek (NF81)				

Wolf Creek

Aquatic Life

Estuary Reservoir River (Sq. Miles) (Acres) (Miles)

Temperature - Total Impaired Size by Water Type: 7.91

Cyala

TMDI

Sources:

Natural Sources Source Unknown

New River Basin

Cause Group Code: N33R-01-BAC Dry Fork

Cause Location: This segment includes Dry Fork south of East River Mountain at the West Virginia state line, downstream to North

Gap (excluding the headwaters).

City / County: Bland Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The AWQM station located at 9-DYF000.07 had 17% exceedance.

Cause Cycle TMDL
Cause First Dev. Water
Assessment Unit / Water Name / Location Desc. Category Cause Name Listed Priority Size
AS-N33R DYF01A12 / Dry Fork / Dry Fork south of East River 5A Escherichia coli (E. coli) 2012 L 5.24

VAS-N33R_DYF01A12 / Dry Fork / Dry Fork south of East River Mountain, the WV state line, downstream to North Gap, excluding headwaters, WQS Section 1 (NE80).

Dry Fork Estuary Reservoir River (Sq. Miles) (Acres) (Miles)

5.24

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

Sources:

Grazing in Riparian or Livestock (Grazing or Shoreline Zones Feeding Operations)

New River Basin

Cause Group Code: N33R-01-TEMP Dry Fork

Cause Location: This segment includes Dry Fork downstream to north Gap, excluding the headwaters.

City / County: Bland Co. Use(s): Aquatic Life

Cause(s) / VA Category: Temperature / 5A

The AWQM station located at 9-DYF000.07 has a 17% exceedance of the temperature.

Cause Cycle TMDL
Cause First Dev. Water
Assessment Unit / Water Name / Location Desc. Category Cause Name Listed Priority Size

VAS-N33R_DYF01A12 / Dry Fork / Dry Fork south of East River 5A Temperature 2012 M 5.24

Mountain, the WV state line, downstream to North Gap, excluding

headwaters, WQS Section 1 (NE80).

Dry Fork

Aquatic Life

Estuary Reservoir River (Sq. Miles) (Acres) (Miles)

Temperature - Total Impaired Size by Water Type: 5.24

Sources:

Grazing in Riparian or Shoreline Zones

Loss of Riparian Habitat

New River Basin

Cause Group Code: N34R-01-BAC Rich Creek

Cause Location: The impaired waters begin just downstream of Peterstown, West Virginia at the mouth of Brush Creek on Rich

Creek and extends to the Rich Creek confluence on the New River (Peterstown, WVA Quad).

City / County: Giles Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The 2002 2.85 mile fecal coliform (FC) bacteria impairment remains. Escherichia coli (E.coli) replaces fecal coliform bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters].

9-RHC000.08 (Rt. 806 Bridge) No additional data collected since the 2016 data window. 2014 and 2016 data windows escherichia coli exceedances occur in 14 of 35 observations. Exceedances range from 275 to 1575 cfu/100 ml. E.coli exceed the 235 cfu/100 ml instantaneous criterion in 14 of 32 samples within the 2012 data window. Exceedances range from 350 to 1010 cfu/100 ml. The 2010 assessment finds E.coli exceed the instantaneous criterion in 10 of 21 samples. Exceedances range from 400 to 1010 cfu/100 ml. E.coli exceed the instantaneous criterion in three of nine samples in 2008 ranging from 400 to 900 cfu/100 ml. Data within the 2006 data window exceed the former FC 400 cfu/100 ml instantaneous criterion in five of nine samples with an exceedance range of 1000 to 2800 cfu/100 ml. The 2004 IR reports FC exceeds the former instantaneous criterion in 10 of 18 samples. Exceeding values range from 500 to 2800 cfu/100 ml.

Cycle TMDI

			Cycle			
	Caus	se	First	Dev.	Water	
Assessment Unit / Water Name / Location Desc.	Categ	ory Cause Name	Listed	Priority	Size	
VAW-N34R_RHC01A00 / Rich Creek / Rich Creek mainstem fro	m 5A	Escherichia coli (E. coli)	2008	Н	2.85	

its mouth on the New River upstream to the Rt. 219 crossing at the Virginia/West Virginia State Line.

Rich Creek

Recreation

Estuary (Sq. Miles)

(Sq. Miles)

Escherichia coli (E. coli) - Total Impaired Size by Water Type:

2.85

Sources:

Municipal (Urbanized High Unspecified Domestic Wet Weather Discharges Wildlife Other than Density Area) Waste (Non-Point Source) Waterfowl

New River Basin

Cause Group Code: N35R-01-BAC **Adair Run**

Cause Location: The Adair Run impairment begins at the Virginia / West Virginia State Line and extends downstream to the Adair

Run confluence with the New River.

City / County: Giles Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 5A

The 2004 303(d) Listed 0.37 mile bacteria impaired waters find the Recreational Use is not supported. Escherichia coli (E.coli) replaces fecal coliform (FC) bacteria as the indicator as per Water Quality Standards [9 VAC 25-260-170. Bacteria; other waters1.

9-ADR000.13 (Rt. 648 Bridge) There are no additional data beyond the 2014 data window. The 2014 assessment finds E.coli exceed the 235 cfu/100 ml WQS instantaneous criterion in five of 32 samples. Values in excess of the criterion range from 325 to 1650 cfu/100 ml. There are no additional data within the 2012 data window. The 2010 assessment finds escherichia coli exceed the 235 cfu/100 ml WQS instantaneous criterion in three of 20 samples. Values in excess of the criterion are 450, 1050 and 1200 cfu/100 ml. The 2004 IR reports fecal coliform exceeds the former 400 cfu/100 ml instantaneous criterion in six of 26 observations. Exceeding values range from 500 to 4200 cfu/100 ml. FC exceeds the former instantaneous criterion in six of 20 observations within the 2006 data window. Exceeding values range from 500 to 4200 cfu/100 ml. FC data within the 2008 data window find four of 14 samples exceeding the former instantaneous criterion.

			Cycle	IMDL	
	Cause		First	Dev.	Water
Assessment Unit / Water Name / Location Desc.	Category	Cause Name	Listed	Priority	Size
VAW-N35R_ADR01A00 / Adair Run / Adair Run mainstem fill mouth on the New River upstream to the Virginia/West Virginia		Escherichia coli (E. coli)	2010	Н	0.37
Line.					

Adair Run Estuary Reservoir River (Sq. Miles) (Acres) (Miles) Recreation Escherichia coli (E. coli) - Total Impaired Size by Water Type: 0.37

Sources:

Unspecified Domestic Wildlife Other than

Waste Waterfowl

New River Basin

Cause Group Code: N36R-01-BAC Bluestone River and Big Branch

Cause Location: This segment extends from Route 460 bridge downstream to the West Virginia political boundary and includes Big Branch from the headwaters downstream to the confluence with the Bluestone River. It also includes Mud Fork, a

Bluestone River tributary at Falls Mills (does not include privately owned reservoir).

City / County: Tazewell Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

Fecal Coliform / 4A

Station 9-BST066.80 had a 56% exceedance of the E.coli water quality standard. The AWQM station located at 9-BST062.47 had a 77% exceedance of the E.coli water quality standard, station 9-BST073.32 had a 37% exceedance. Stations 9-BIG000.12 had a 88% exceedance and 9-MFK000.11 had a 36% of the E. coli water quality standard.

Assessment Unit / Water Name / Location Desc.	Catago	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N36R_BST04A02 / Bluestone River / From Wright's Valley Creek confluence downstream to N37 at the Big Branch confluence below Falls Mills, section 1g, u (NE86).	4A	Escherichia coli (E. coli		2004	L	6.23
VAS-N36R_BST04B02 / Bluestone River / From PWS intake for Town of Bluefield, downstream to Wright's Valley Creek confluence section 1g, u (NE86).		Escherichia coli (E. coli)	2006	L	1.72
VAS-N36R_BST05A02 / Bluestone River / From Town of Bluefie PWS intake, upstream to Rt. 460 bridge near Shannandale, WQS Section 1h, u (NE86).	ld 4A	Escherichia coli (E. coli	1	2006	L	5.05
VAS-N36R_MFK01A06 / Mud Fork / Bluestone tributary at Falls Mills, north of Stony Ridge upstream to SR 608 bridge, WQS Section 1g. Does not include privately owned reservoir (NE87).	4A on	Escherichia coli (E. coli	•	2018	L	2.98
VAS-N37R_BIG01A10 / Big Branch / Bluestone tributary south o Abbs Valley Ridge, from headwaters in WQS Section 1g, parallel S 698 (NE 88).		Escherichia coli (E. coli	1	2010	L	3.33
VAS-N37R_BST01A96 / Bluestone River / Mainstem from Big Branch confluence downstream to WV state line near Yards in WQ Section 1g, u (NE88).	4A S	Escherichia coli (E. coli	1	2006	L	0.62
Bluestone River and Big Branch Recreation Escherichia coli (E. coli) - Total	Impaired	I Size by Water Type:	Estuary (Sq. Miles		servoir .cres)	River (Miles)
		. 0.20 2)				10.00
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N36R_BST04B02 / Bluestone River / From PWS intake for Town of Bluefield, downstream to Wright's Valley Creek confluence section 1g, u (NE86).		Fecal Coliform		2002	L	1.72
VAS-N37R_BST01A96 / Bluestone River / Mainstem from Big Branch confluence downstream to WV state line near Yards in WQ Section 1g, u (NE88).	s 4A	Fecal Coliform		2002	L	0.62
Bluestone River and Big Branch			Estuary (Sq. Miles		servoir	River (Miles)
Recreation Fecal Coliform - Total	Impaired	Size by Water Type:	(Oq. Miles	, (A	10163)	2.34
- Total Comolin - Total	paniec	. C.LO by Traidi Type.				2.04

New River Basin

Sources:

Rural (Residential Areas)

Sewage Discharges in Unsewered Areas Silviculture Activities

Source Unknown

New River Basin

Cause Group Code: N36R-01-BEN Bluestone River

Cause Location: This segment extends from the Wright's Valley Creek confluence downstream to the West Virginia political

boundary.

City / County: Tazewell Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

Biological station 9-BST069.21 was impaired based on VSCI scores of 49 and 53 in 2016. Station 9-BST069.07 was impaired based on VSCI scores of 58.2 and 56.3 in 2017.

Benthic Macroinvertebrates Bioassessments - Total Im	paire	d Size by Water Type:				6.85
Bluestone River Aquatic Life			uary Miles)		ervoir cres)	River (Miles)
VAS-N37R_BST01A96 / Bluestone River / Mainstem from Big Branch confluence downstream to WV state line near Yards in WQS Section 1g, u (NE88).	4A	Benthic Macroinvertebrates Bioassessments	20	02	L	0.62
VAS-N36R_BST04A02 / Bluestone River / From Wright's Valley Creek confluence downstream to N37 at the Big Branch confluence below Falls Mills, section 1g, u (NE86).	4A	Benthic Macroinvertebrates Bioassessments	20	02	L	6.23
_	Cause atego	e ry Cause Name	Cyo Fir List	st	Dev. Priority	Water Size

Sources:

Crop Production (Crop Land Silviculture Activities or Dry Land)

Unrestricted Cattle Access

New River Basin

Cause Group Code: N36R-01-CDANE **Bluestone River**

Cause Location: This segment includes the mainstem from the confluence with Big Branch downstream to West Virginia political

boundary; may be found on the Bramwell quad sheet.

City / County: Tazewell Co. Use(s): Fish Consumption

Cause(s) / VA Category: Chlordane / 5A

The fish tissue and sediment sampling stations at 9-BST069.46 and 9-BST066.94 had total chlordane levels detected in the

sediment in 2002 above DEQ's screening value.

Cycle **TMDL** Dev. Cause First Water Assessment Unit / Water Name / Location Desc. Category Cause Name Listed **Priority** Size Chlordane 2004 0.62 VAS-N37R BST01A96 / Bluestone River / Mainstern from Big

Branch confluence downstream to WV state line near Yards in WQS

Section 1g, u (NE88).

Bluestone River **Estuary** Reservoir River (Sq. Miles) (Acres) (Miles) **Fish Consumption**

Chlordane - Total Impaired Size by Water Type:

0.62

Sources:

Source Unknown

New River Basin

Cause Group Code: N36R-01-PCB Bluestone River

Cause Location: This segment begins at the Route 460 bridge downstream to the West Virginia political boundary. It also includes a segment of Beaverpond Creek that flows from West Virginia into Virginia, sometimes under city buildings and streets and into the Bluestone River and Brush Fork from the west Virginia state line to the confluence with the

Bluestone River in Falls Mills.

City / County: Tazewell Co. Use(s): Fish Consumption

Cause(s) / VA Category: PCBs in Fish Tissue / 5A

Polychlorinated Biphenyls (PCBs) / 5A

In April 2004 a Special Study was conducted by DEQ and USGS. An SPMD deployed at station 9-BPB000.02 indicated Total PCBs in the water column at 3700pg/l and 1300pg/l in 2005. SPMDs deployed at stations 9-BST066.18, 9-BST068.98 and 9-BST072.65 indicated PCB values of 1800 pg/l, 800 pg/l and 230 pg/l. Fish tissue and sediment stations 9-BST0666.94 and 9-BST069.46 found PCB in exceedance of DEQ's screening values in white suckers. Station 9-BST069.46 also had sediment samples that exceeded the ER_M for PCBs. SPMD sampling in 2004 indicated PCB was 3500 pg/l at station 9-BFK003.14.

	Cause	4		Cycle First	TMDL Dev.	Water
Assessment Unit / Water Name / Location Desc.		ry Cause Name		Listed	Priority	Size
VAS-N36R_BST04A02 / Bluestone River / From Wright's Valley Creek confluence downstream to N37 at the Big Branch confluence below Falls Mills, section 1g, u (NE86).	5A	PCBs in Fish Tissue		2002	Н	6.23
VAS-N36R_BST04B02 / Bluestone River / From PWS intake for Town of Bluefield, downstream to Wright's Valley Creek confluence, section 1g, u (NE86).	5A	PCBs in Fish Tissue		2002	Н	1.72
VAS-N36R_BST05A02 / Bluestone River / From Town of Bluefield PWS intake, upstream to Rt. 460 bridge near Shannandale, WQS Section 1h, u (NE86).	5A	PCBs in Fish Tissue		2002	Н	5.05
VAS-N37R_BST01A96 / Bluestone River / Mainstem from Big Branch confluence downstream to WV state line near Yards in WQS Section 1g, u (NE88).	5A S	PCBs in Fish Tissue		2002	Н	0.62
Bluestone River			Estuary	Res	servoir	River
Fish Consumption			(Sq. Miles) (A	cres)	(Miles)
PCBs in Fish Tissue - Total Ir	npaired	Size by Water Type:				13.62
Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name		Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N36R_BFK01A06 / Brush Fork / Bluestone tributary from W\state line downstream to Bluestone River at Falls Mills parallel to SR 643, WQS Section 1g (NE86).		Polychlorinated Bipheny	rls (PCBs)	2010	Н	1.48
VAS-N36R_BPB01A06 / Beaverpond Creek / Bluestone tributary from WV state line, sometimes under town buildings and streets in Bluefield, downstream to Bluestone confluence, WQS Section 1g (NE86).	5A	Polychlorinated Bipheny	rls (PCBs)	2012	Н	2.99
Bluestone River			Estuary		servoir	River
Fish Consumption			(Sq. Miles) (A	cres)	(Miles)
Polychlorinated Biphenyls (PCBs) - Total Ir	npaired	Size by Water Type:				4.47

New River Basin

Sources:

Illegal Dumps or Other Inappropriate Waste Disposal Source Unknown

New River Basin

Cause Group Code: N37R-01-BAC **Laurel Fork**

Cause Location: Laurel Fork mainstem from the Curran Branch confluence at Boissevain to the WV state line east of Pocahontas.

City / County: Tazewell Co.

Use(s): Recreation

Cause(s) / VA Category: Escherichia coli (E. coli) / 4A

The AWQM station located at 9-LRR001.39 had a 83% exceedance of the E.coli water quality standard.

Cycle **TMDL** First Cause Dev. Water Listed Priority Category Cause Name Size Assessment Unit / Water Name / Location Desc. Escherichia coli (E. coli) 2006 4.70 VAS-N37R_LRR01A94 / Laurel Fork / Laurel Fork mainstem from the Curran Branch confluence at Boissevain, to WV state line east of

Pocahontas, WQS Section 1 (NE88)

Laurel Fork **Estuary** Reservoir River (Sq. Miles) (Acres) (Miles) Recreation

> Escherichia coli (E. coli) - Total Impaired Size by Water Type: 4.70

Sources:

Sanitary Sewer Overflows Septage Disposal (Collection System Failures)

New River Basin

Cause Group Code: N37R-01-BEN Laurel Fork

Cause Location: Laurel Fork mainstem from the Curran Branch confluence at Boissevain to the WV state line east of Pocahontas.

City / County: Tazewell Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

The biological station at 9-LRR001.39 found that the segment was impaired based on the VSCI.

Cycle **TMDL** First Cause Dev. Water **Priority** Category Cause Name Listed Size Assessment Unit / Water Name / Location Desc. 4.70 VAS-N37R_LRR01A94 / Laurel Fork / Laurel Fork mainstem from Benthic Macroinvertebrates 1996 Bioassessments the Curran Branch confluence at Boissevain, to WV state line east of Pocahontas, WQS Section 1 (NE88)

Laurel Fork Estuary Reservoir River
Aquatic Life (Sq. Miles) (Acres) (Miles)

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type: 4.70

Sources:

Impacts from Abandoned
Mine Lands (Inactive)

Silviculture Activities

New River Basin

Cause Group Code: N37R-02-BEN Laurel Fork

Cause Location: Upstream of the Curran Branch confluence at Boissevain to headwaters on Yokel Ridge (parallel to the West

Virginia state line).

City / County: Tazewell Co.

Use(s): Aquatic Life

Cause(s) / VA Category: Benthic Macroinvertebrates Bioassessments / 4A

Probabilistic monitoring station at 9-LRR012.30 was impaired based on VSCI scores of 38.6 and 44.4 in 2014, 54.7 and 56.3 in

2016, and 51.31 and 50.38 in 2018.

Assessment Unit / Water Name / Location Desc.	Cause Catego	e ry Cause Name	Cycle First Listed	TMDL Dev. Priority	Water Size
VAS-N37R_LRR02A02 / Laurel Fork / Upstream of the Curran Branch confluence at Boissevain to headwaters on Yokel Ridge (parallel WV state line), WQS Section 1 (NE88).	4A	Benthic Macroinvertebrates Bioassessments	2014	L	8.30

Laurel Fork

Aquatic Life

Estuary Reservoir River
(Sq. Miles) (Acres) (Miles)

Benthic Macroinvertebrates Bioassessments - Total Impaired Size by Water Type: 8.30

Sources:

Source Unknown